

Figure 1BF V

B-F V

GGC CCC GGG ATG CCG CGG TTC GTG ATC GTC GGG TAC GTG GAC GAC AAA ATC TTC GGT
ACC TAC AAC AGT AAG AGC AGG ACT GCA CAG CCT ATC GTG GAG ATG CTG CCG CAG GAG
GAC CAG GAG CAC TGG GAC ACG CAG ACC CAG AAG GCG CAG GGC GGT GAG CCG GAT TTT
GAC TGG AAC CTG AAC AGG CTG CCG GAA CGC TAC AAC AAA AGT AAA GGT GAG CGT GGG
GGA AGC TGC AGC GCG ATG CGT CTG GGA CAG GAG CTC TGT GTG CCG AGG GTG TCC GCC
AGC CCC ACT GAG GTG TGG CCG TGC CCC ACG CCC AGC TGT GCT GGG CCG TCC ATG TGT
GGT GGC ACT GTC CCT GGG CCG CCC TGC TCC TGC GCC CAC CCA CCC CAC CCC AGC CTC
ATG GCA CTC GCG GTG CCC CAC AGC CCT AGA AGC CTC TCA CCT ATT ACT CTG GCT GTG
CCT CAG GGT CTC ACA CGA TGC AGA TGA TGT TTG GCT GTG ACA TCC TGG AGG ACG GCA
GCA TCC GAG GGT ACG ATC AGT ATG CAT TTG ATG GGA GGG ACT TCC TTG CCT TTG ATA
TGG ACA CGA TGA CGT TCA CCG CGG CGG ATC CAG TGG CTG AAA TCA CCA AGA GGA GAT
GGG AGA CAG AAG GGA CGT ATG CTG AGA GAT GGA AGC ATG AGC TGG GGA CTG TCT GTG
TTC AGA ACT TGA GGA GAT ACC TGG AGC ATG GGA AGG CAG CGC TGA AAA GGA GAG GTG
AGG ATG GGA GGG GGA CGT GGG GCT GGG CTG GGT GTG GGG CAG AGG CTC AGT GTG GGG
TGC TCA GCC CGG CCC ACA ACG TCA CCC ACC TGC AGT GCA GCC CGA GGT GCG AGT GTG
GGG GAA GGA GGC CGA TGG GAT CCT GAC CTT GTC CTG CCA CGC TCA CGG CTT CTA CCC
GCG GCC CAT CAC CAT CAG CTG GAT GAA GGA CGG CAT GGT CCG GGA CCA GGA GAC CCG
CTG GGG GGG CAT CGT GCC CAA CAG CGA TGG CAC CTA CCA CGC CTC GGC TGC CAT TGA
TGT GCT GCC GGA GGA TGG GGA CAA GTA TTG GTG CCG CGT GGA GCA CGC CAG CCT GCC
CCA GCC TGG TCT CTT CTC ATG GGG TGA GCT GGC AGC GTG GGG CAC GTG GGG TTG GGA
TTC GCA GGC TGC CCC TTC CTT TAC TGA CAA CGG CGC TCT CCT CCA GAG CCG CAG CCC
AAC CTG ATT CCC ATT GTG GCA GGG GCG GTC GTT GCC ATC GTG GCT GTC ATC GCT GCG
GTC GTT GGA TT

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Figure 2B-FVI

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ATC TAC GAC AGT AAG AGC AGG ACT GCA CAG CCC ATC GTG GAG ATG CTG CCG GAG GAG
GAC CAG GAG CAC TGG GAC GCG CAG ACC CAG AAG GCC CAG GGC GGT GAG CCG GAT TTT
GAC TGG TTC CTG AGC AGG CTG CCG GAA CGC TAC AAC AAA AGT GGA GGT GAG TGT GGG
GGA AGC TGC AGC GCG ATG CGT CTG GGA CAG GAG CTC TGT GTG CCG AGG GTG TCC GCC
AGC CCC ACT GAG GTG TGG CCA TGC CCC ACG CCC AGC TGT GCT GGG CCG TCC ATG TGT
GGT GGC ACT GTC TCT GGG CTG CCC TGC TCC TGC GCC CAC CCA CCC CAC CCC AGC CTC
ATG GCA CTC GCG GTG CCC CAC AGC CCA AGA AGC CTC TCA CCT ATC ACT CTG ACT GTG
CCT CAG GGT CTC ACA CGA TGC AGA TGA TGA TCG GCT GTG ACA TCC TGG AGG ACG GCA
GCA TCC GAG GGT ACG ATC AGT ATG CAT TTG ATG GGA GGG ACT TCC TTG CCT TTG ATA
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GGG AGA CAG AAG GGA CGT ATG CTG AGA GAT GGA AGC ATG AGC TGG GGA CTG TCT GCG
TTC AGA ACT TGA GGA GAT ACC TGG AGC ATG GGA AGG CCG CAG TGA AAA GGA GAG GTG
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TGC TCA GCC CCG CCC ACA ACA TCA ACC ACC TGC AGT GCA GCC CGA GGT GCG AGT GTG
GGG GAA GGA GGC CGA TGG GAT CCT GAC CTT GTC CTG CCA CGC TCA CCG CTT CTA CCC
GCG GCG CAT CGC CAT CAG CTG GAT GAA GGA CAG CAT GGT CCA GGA CCA GGA GAC CCG
CTG GGG GGG CAT CGT GCC CAA TAG GGA TGG CAC TTA CCA CAC TTC GGC TGC CAT TGA
TGT GCT GCC GGA GGA TAG GGA CAA GTA TCG GTG CCG CGT GGA GCA CGC CAG CCT GCC
CCA GCC TGG CCT CTT CTC TTG GGG TAA GCC TGG CAG CGT GGG ATG TGT GGA GTT GGG
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TTT GAT TGC ATT

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 GCG GAG AGC AAT GGC TGC GGG GCT GCG GCT GCT GCT GGC GGG TGA GAC CCG
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 GCT GCG CGC TGC TGG AGG GGG TGG GGC GCG GGG GAG GGC TGC CGG GGG GGG
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 AGA GTC CCC CAA ACC CCC ATG ACC CCC CCC ACG ACC CCT GGT TCC CAT TAC
 CCT CTC ACG TCC CCC AGT GTC CCC AAG ATT CCC ATT ACT CCC CGT ATC CCC
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 ATG TCC CCA AAA TCC CCA TTA TCT CCC ACC TCT CCA AAG TCC CCA AGA TCC
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 ACC GGG TGG TAT TCC TAT GAG GAC TAT AGG ATG GGG TGG CAT EAT CCC ATA
 GTT CAC CTG TAG GTT TAT AGG GGG GGA TGA GCC CTA TAC AGC GTA TGG GCT
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 AGT GGG AGC AGT GGG AGG AAG TCC TGG GTT GGT GAG GCA GAT GAG TGG CAC
 CTG GGG ACA TCT GGG TGC CAT CCC TTG TGG ACA TCT GGG TGA CAC TGC ATT
 GCC TTG GGT GAC ATT GGG ATC CTC AGG TCA CTG CAG

B-FI

GT CGA CCGGAT CTGGATAGGT CGT CAGT CAT CCTAATTAAGGAGGGGA CAA CAGTGAATGGG
 GAGGAG CCGATGACT CAGG CTGGGAGTGGTGAT CCCAGAGGTTT CCT CTG CTGT CAGTGAC
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 ATTT CCGT CT CCCCCCCCCCCCCCCCCCCCCCG CCGAT CCGAGTGCT CTG CAACT CCT CAC
 CCTG CCGA CACAACTTTG CG CACT CCACT CCGT CACT CCGC CCGT CCCCCAG CT CT C
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2017-10-03 14:00

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FEUILLE DE REMPLACEMENT (REGLE 20)

FEUILLE DE REMPLACEMENT (REGLE 26)

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 GCCCACGGTGTGATCCCCCCCCAGCAGCAACACAGGGAGCTCCCATGGGGT
 TGAGTTTGGGTTCTCAGGGTTTGTCTGTCCCCCATTTCCCACCACCCC
 TTTGGGTTCTCACCAGCAGGAATTTGCTGTGCGGGCTGCTGGAATTTGCCC
 TCCATCTCCCAGATCAGGGTGTCAAGGTGGGACATCTCCTCCATCACCTT
 CGTCACCGCATCCTCCTGTACTTTGGTGACGGCTCTGTCCAGGTCTGCCA
 GCTGGACCAGCAGGAAGCGCTCCTTCTCCTTCAGAAATCGCTGCAACTGC
 TCGAATTCACACACTATCCTCTTCCCTTCTTCTTGGTTTTCTCCTGTTG
 GGATGAGGGAGAAAGCCAATGGGGTGGAATAGAGGCAGGAAGACCCCCC
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 GGATGTAACACCAATGCCAATGGGAGCACAACACTAATGCCAATGGGAAT
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 CACCAGTGCCAGTGGGAATTTATCACCAGTGCCAATGGGAACCTTAACATC
 AAAAAGCCAAAGATCATCTTGCTGGGCATTTGGGAGCAGCAGGAATTTTT
 CAGGAGTTTTATCCCAAAAGCAAAACCAAGGAGGGGGTAGGAGATGAGC
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 TTTAACACAGGGGTTCTGCGTAACCCCAGCTGATAAACACAGCCTTAGC
 GCTTTCCCAGCCCAGCTGCGAGCCAAAATGCATGATCTGCCCCCAAAT
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 GCTTCTGTGATGGGAGAGGCTTTGCTAAAGCCTGGAATCCTCTGCAAG
 GTGCAGAGCTGGGCAGAGGGAAGCTCTGTGAGCACGGTGTGCTGCTCTGG
 AGCTCTGTGCAAGCTGGGAGTATTTTGCAGAGAGAAAAGAGGGGAGAAAG
 GAAGGAAAAACAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG
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 AACCCACAGAACCTGTTGTTTTAGCCCTTTGATGGGGTTGGGGTTTTCC
 CTTCTGTTCCTTCCAGTCTGGGGTAGAGCTATGGGATGGCTGCGTTGA
 GCCTGCAGGTCTGCTCCTGGTGGCACCCCTTGGCAGGGCGTGCTGGGAGCT
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 GAAGTGCATTACAGCGTAGGTTGGCACGGGGAGACGTGAGGGAGTGGTAG
 GCCTCCAGCGCCAGACACCCCTCTTTGGGGCTGAAGCTCATGGGTCCCTT
 CCTCTTCATCGAAGC **EFFICACE DE REMPLACEMENT (REGLE 26)** CCCCTGTC
 CCACCTCCACCTCCCAGAAATGCCTCCCCGAGGTGAAGCCCTGGCAGCCC

ACAAAGATGAGCCCCCTTCATCAGCATCAAAAAATGCCACCGTCCCTCC
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 CCTGATCGACAAGACCGGCTTCCATCCGAGTACGTGCTCGCTCGATGCGA
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 GGAGGGCGGGGGGGTGTAAAGAGGATCGCGGCGCTCTTCTGCGTGCCCG
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 CATCCTCGGCACCGTCACCCTGGATGCTGTAGGCATAGGCTTGTTATGC
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 GCTTTGTGACATCCAGCGCCGACATTCAGCAGCGTTTTTCAGCGCGTTT

DM

GCACAAGGAA ATGCAAAGGG GCATCACTAG GGGACATGGC ACGGGGCATT
51 CTAGGGAGCA TTGCATGGGG ACATTGCAAA GGAAATGCA AGGGACATTG
101 CATGGGGACA TTGCAAACAA ATTGAGTGGG AGATTGCACC GGGATGTTGC
151 ATGGGGACAT TGCATGGAAT GTCCCACCA CCACCCCTGCA GGGTGACACT
201 GGGACCATCC CCAGCTCTGA CCATCCCCC TTTGCTGCAG CACCACCCCA
251 GGTCCGCATC GTCCCCATCC CCATCTCAA CGACCCCGAC ACCGTCCACC
301 TCATCTGCCA TGTTTGGGGC TTCTACCCAC CCGCAGTGAC CATCCAGTGG
351 CTGCACAACG GCCTCGTGGT GGCCTCAGGT GACACCAAAC TGCTGCCCCA
401 CGGGGGACTG GACCTACAGG ACACAGGTGG CCTGAGGGC CAGCATTGCA
451 GCAGGGAGCA CTAAAACATG TTCAGTGTGG CAATCCAGC TTGGAGCAGC
501 CGCTGCAGGA GGATTGGAGT GAGTTTGGGG ATGGGGATGT GGCACCCACA
551 CCCCACAGTC CCCCACGGTT CATTGTGCCC CACGCTGTCC CCACAGGTCC
601 CAATTTGTCC CCGGCGATGA TGGTGAAGGT GGCAGTGGCG GCCATGGCGC
651 TGACGTTGGG GTTGGTGGCA CTCAGCGCCG GGGTTTTTCAG CTTCTGTGAG
701 CGGCCACGGG GTGAGGGATG GGGATGTGGT GCTGGGGACA TGTGTGACAC
751 CGAGGGTCTG GTGTCCAGTG TGGGGTGTAC CTCCTCATTC ATCATCTTCT
801 GTGTGGCAGC TCCTGGCGCT GGTCCCAGTA CCCCCTCCTG ATGCGGGTTC
851 TCACTCCAAT CCTGGTCCCC AAAATGATCC CGGTCCGAGT TCTGGTCCCC
901 ATCCCAGTCC TGGACCCCAT CCCAGTCCTG GTCCCCATTC TGGTCTTGGT
951 CCTGGTCTCTG GTTCTGTCTC TGGTCCCTAT CCCTGACTCT GGTCCCGGTC
1001 CCCATCCCGA TGCCAGTCCC AGTCCTGGTC CCCATCCTGG TCCTGCTCCT
1051 TGGTTTGGGG ACCTCAATGA CTGGAACTCC CATGTCCCAA CATGGGGACC
1101 CACAGTTTGG GGTGAGGGGC TCTCACCCCC CAATAAAACC ATCTGCAGCC
1151 CCAACCTCGC TCCAATTCTT CGTTCCCACG TTGGGTGGGT CGGGCTCCCA
1201 GTGCTCCAG CCGTNTATGT CCCGTAAGCG TCGGCTCCAC TGCATAAAAA
1251 GAAAAAAAA AAA

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Figure 7

Séquence Génomique TAP1

(Du début de l'exon 2 à l'extrémité 3')

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GGC GAG ATG GCC GTG CCC TAC ATG GGG CGA GCC AGC GAC TGG GTG GCC CGC GAG GAC AAG CTG GCA
G   E   M   A   V   P   Y   Y   M   G   R   A   S   D   W   V   A   R   E   D   K   L   A

GCC ATC CTG CCC ATG GTG CTG GGC CTC AGC AG GTACTGGCATAGGGGGGACGGGTGCGGGCAGGGGGCAGCGCG
I   L   P   A   M   V   L   L   G   L   S   S

GGACCCCTGACACCCCACTGCCGTACAG C GCT GTT ACT GAG CTG GTG TGT GAT GTG ACC TTC GTG GGG ACA
A   V   T   E   L   V   C   D   V   T   F   V   G   T

CTG AGC CGC ACG CAA AGC CGC CTC CAG CGC CGC GTC TTC GCC GTC CTC GTG CGG CAG AGC ATC ACC GAG
L   S   R   T   Q   S   R   L   Q   R   R   V   F   A   A   V   L   R   Q   S   I   T   E

CTG CGC GCC GAT GGG GCC G GTGAGGGGACCGGGCTGGGAGGGGACACCGGGGATPAGGGACAGGGGTGGCCTGACGGCGCTG
L   R   A   D   G   A

TCACCCCGGCAG GG GAT GTG GCC ATG CGG GTG ACG CGG GAT GCG GAG GAC GTG CGC GAG GCG CTG GGC AAG
G   D   V   A   M   R   V   T   R   D   A   E   D   V   R   E   A   L   G   K

GCG CTG AGC CTC CTG TGG TAT CTG GCA CGC GGC CTC TGC CTC TTT GCA ACC ATG GCC TGG CTG TCC
A   L   S   L   L   L   W   Y   L   A   R   G   L   C   L   F   A   T   M   A   W   L   S

CCG CGC ATG GCG CTG CTC ACC GCG CTG GCG CTG CCA CTG CTG GCA CTG CCC AGG GCT GTG GGG CAC
P   R   M   A   L   L   T   A   L   A   L   P   L   L   L   A   L   P   R   A   V   G   H

TTC CGG CAG GTATGGGCTGCTGTGTCACCTCCATGTGCCCTTTGGTCCCTCCATGTGCCTCTGGTCCCTCCATGTGCCAGTGTC
F   R   Q

```

ACCACCATGTACTCATTTGCCCTATCCATGTGCCACATGTCCCCCTCCATGTACCCACCATCCTCCTGCTGTGTGCCCTCTGTGTGACCGGC
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 A L A
 CCA CAG ATG CAG AAG GCG CAG GCC CGG GCC AGC GAG GTG GCA GTG GAG ACC TTC CAG GCC ATG GCC ACT
 P Q M Q K A Q A R A S E V A V E T F Q A M A T
 GTG CGC AGC TTT GCC AAT GAG GAT GGG GCA GCT GCA CAC TAC CGG CAG CGC CTG CAG CAG AGC CAC CGC
 V R S F A N E D G A A H Y R Q R L Q Q S H R
 CTG GAG AAA AAG GAT GTG GCC CTC TAC ACT GCC TCT CTC TGG ACC AGT GGT GTATGGGATGGGGTGGCTCAAT
 L E K K D V A L Y T A S L W T S G
 AGCATGGGACGTGATGGGATGGGGCTGGGGATGTGGGGACATGATAGCATAGGACTGGGGGGCATGGGGACATGGTGGGATAGGGCTGG
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 F S A L A L A L K M G I L Y Y G
 GGG CAG CTG GTG GCC GCG ACC GTC AGC ACT GGG GAC CTC GTC ACC TTC CTC CTC TAC CAG ATA CAG
 G Q L V A A G T V S T G D L V T F L L Y Q I Q

FIGURE 7 - SUITE 1

Figure 7

TTC ACT GAT GTC CTG GAG GTGAGCCTGAGAGGATGCCCATATCCGCATGTCCCCATGTCTCCCTGCCACAGTCACAGTGTGTCA
 F T D V L E

CAGTGTCCCTGTCTTCTACATCCTCCCTGTGTCTCCAGGCCAATCCAGTGTGTCCCCATGCCATCCATGTCCCTGTGTCCCCACG
 TCTAACCTGTGTGACCCCTGCCACATCCCTGTGTGCACCTCCTCTGTCCCCATACCTACCCCGTCTGTCCCTATGCCAATCCCACTGTGT
 CCTGTGGTCCCCATGTACACCATGCCACATGCCCTTCCCCCTCTGTGCCACGTCACTACGCTGTCCCCAG GTC CTG CTC GAC TAC
 V L L D Y

TTC CCC ACA CTG ATG AAG GCT GTG GGC TCT TCG GAA AAA ATC TTT GAG TTC CTG GAC CGG GAG CCA CAG
 F P T L M K A V G S S E K I F E F L D R E P Q

GTC TCA CCC TCA GGG ACA ATG GCA CCC GCT GAC CTG CAG GGC CAC CTC CAG CTG GAG GAT GTC TGG TTC
 V S P S G T M A P A D L Q G H L Q L E D V W F

TCC TAC CCT GGG CGC CAG GAA ACC CGT CCT CAA GTGGCACAGAGACACAGCCAGGGGACACGGGGGTGTGGTGGGACA
 S Y P G R Q E T R P Q

GGTGCACAGGTGTGGAGCACAGTGGGGTGTATCAGGGACATGGATGTATGGACAGGGGTGTGAGGATATGAAACAAGGACATACATGGAGG
 GGGTGGTATGGGGACACTGGAGAGGGACATGAGATCATGGTATTGAGGGCGCGGGGACATGGCACATGGTGGGTGTGGCACATGGGACAT
 GATGAGTGACACAGAGACATGGTGGGAGGGGATGGGAATGTAGAGGCCGTGGTA

GGG GTA TCA CTG GAG CTG CGC CCC GGG GAG GTG GCA CTG GGA CCC CCG GGC GCA GGG AAG AGC
 G V S L E L R P G E V L A L L G P P G A G K S

ACT CTG GTG GCC CTC GTG TCC CGC CTG CAC CAG CCC ACG GCC GGG CGC CTG CTG GAT GGC CAC CCC
 T L V A L V S R L H Q P T A G R L L L D G H P

CTC CCC GCC TAC CAG CAC TCC TAC CTG TGC CGC CAG GTGAGCAGCCACATGTCCCCATGGCTCCTGGTGTCCCCCTG
 L P A Y Q H S Y L C R Q

FIGURE 7 - SUITE 2

FIGURE 7 - SUITE 3

FIGURE 7 - SUITE 4

CGG GCA CAA CGA GTG GTG TTG GAG GGG GGA GAG GTG CGG CAG GAG GGA CCC CCC CAC GAG GTG GTG
P A Q R V V V L E G G E V R Q E G P P Q E V V

CGC CCC GTC AGC CTT NTT GCG GGA CTG GGG ACA ACA AGG GAG CAC CGG GGG AGG GGG ACA GAG GGA TAG
R P V S L ? A G L G T T E H R G R G T E G *

CGGGAGTTTGGATGGGGAGGGCGGGGTGGTGGGATGTGGGATGGGGACACTGCGCGTTGGGGACACTGAGGGTGGAGGTGGGGACAC
CGGGGCAGACCAACAGGGACCACACAGCTGTCCGTGGGCACATGGATGCCGAGCCGGCGCGTCCGGTACCGTGTGTACGACACACA
ACGGCCACAGCATTGGACTGCAGTGGCAGTGGCCACCGGCCCGGGGTGGGACACAGAACTGGGAATAAAGCCGCATGTTTGT

Figure 8

TAP2G

-213
CGCCATACATNTGCGCCTGTCTATGCACGGTGNATATGGCCGACCTGGCCNVCATGTTGGCCCTGGCCCANITCTTCCCAGCACTGGCCCCA
TTGGCGTGGTGGC -107
-106
TCCTGGCGGAGGCCGGGCTCCTGGTGTCTGGGGGGGGGGGGGCGACCTGCTGGCCCCCAGGGGACCCCGTGGGGCTGCAGTGCTGCT
AGCATGGCCCCGCC 0
t
+1 ATCTTCCTGACCCCTACGGGGCTATGATGCTGCTGAGCTCCCCCGGTGCTGCTGGCC ATG GCA ACG CCG TC
TGG CTG GTG CTG ACC CAC +93 5'DT
W L V L T H M A T P S
+94 GGG ACA GCT GTG GCA TTG CTC ACC TGG AGC CTC CTG GTC CCC ACT GTG GCC ACT GGG
GCA AAG GAG GCA AAG GCC TGG +174
G T A V V A L L T W S L L V P T V A T G
A K E A K A W
+175 GTG CCC CTG AGG CGG CTG CTG GCC CTC TGG CCC GAG TGG CCG TTC CTT GGC TGT GCC
TTC CTC TTC GCA TTG GCT +255
V P L R L L A L A W P E W P F L G C A
F L F L A L A
+256 GCA CTG GGT GAG ACC TCA TTG CCC TAC TGC ACC GGG AGG GCT GTG GAT GTC CTC CGC CAG
GGG GAC GGC CTC GCC GCC TTC +336
A L G E T S L P Y C T G R A V D V L R Q
G D G L A A F
+337 ACC GCT GCT GTC GGC CTC ATG TGC CTG GCC TCT GCC AGC AG
GTAGGGACCCACATCCCTCCACAAACCCCATCCACCTCTGGTGGTCT +429
T A A V G L M C L A S A S S

+430
GGTGGGTTGGGGGTCTCTGTCCATATCTGGGGGTCTATCTGATGGGTCTGTGGGCACCTCCACTGACCCCTTGTGATGTCTGAAGGGTCTCTG
GCTCTCCATGACCC +536

+537 CTGATGGGTTTGGAGTCCGCCCCCAATTCCTTCCAG C TCG CTG TTT GCC GGC TGC CGC GGT GGC CT
TTC ACC TTC ATC AGG TTC +624
S L F A G C R G G L

F T F I R F

+625 CGC TTC GTC TTG CGC ACC CGC GAC CAG CTC TTC TCC AGC CTG GTC TAC CGG GAC CTC GCC
TTC TTC CAG AAC ACC ACA GCA +705
R F V L R T R D Q L F S S L V Y R D L A

F F Q N T T A

+706 G GTACAGACTGGGGGCACCTTTTGTCCCTGTCCCCACACCCTACCCCGTACTCACTCCACAG CT GAG
TTG GCC TCC CGG CTG ACC ACC +828
A E

L A S R L T T

+829 GAT GTG ACG CTG GCG AGC AAC GTG TTG GCA CTC AAT ATC AAC GTC ATG CTG AGG AAC CTG
GGG CAG GTG CTG GGG CTC TGC +909
D V T L A S N V L A L N I N V M L R N L

G Q V L G L C

+910 GCC TTC ATG CTG GGG CTG TCC CCG CGC CTG ACA ATG CTG GCA CTG CTC GAA GTG CCG CTC
GCC GTC ACC GCA CGG AAA GTC +990
A F M L G L S P R L T M L A L L E V P L

A V T A R K V

FIGURE 8 - SUITE 1

Figure 8

+991 TAT GAC ACC CGG CAC CAG
GTGATAGCAGGGATGGGATGGTAGGGTTGGGGTGCACAGGGATGGAGGCAATGGCAATGGGATGGGAACAGTGGGAGTGGGGAT +1091
Y D T R H Q

+1092
AGTCAGGTGGGGATTTGGGGTTCAGGGTGGCAGGGATGAGGGCAGCTGCAATGGGATGGGAACAGTGGGAATGGGAGAGCAGGATGGGGA
CATGGGTCCACACACA +1198

+1199
GCAAGGATGAGAGGATGGGAAGAGTGGAGCAGGAATGGAACTGGGATGGCGAGTACTTGGCCATCCCATGGGTGCTGACACCCACTGTCC
CCCCAG ATG CTG +1302

M L

+1303 CAG CGG GCC GTG CTG GAT GCA GCA GCC GAC ACC GGA GCG GCA GTG CAG GAG TCC ATC TCT
TCC ATT GAG ATG GTA CGG GTC +1383
Q R A V L D A A A D T G A A V Q E S I S

S I E T V R V

+1384 TTC AAT GGC GAG GAG GAG GAG GAG CAC CGC TAC AGC CAG GTG CTG GAC AGG ACC CTA CGG
CTG CGG GAC CAG CGG GAC ACA +1464
F N G E E E E H R Y S Q V L D R T L R

L R D Q R D T

+1465 GAG AGG GCC AAT TTT CTC CTC ATC CAG CGG
GTGAGGCTGACACGAGGGGACACCCCTGGTGTCTGGGTGGATCGGGACATCCCCCGCTGAGCCCCAT +1561
E R A I F L L I Q R

+1561 CCCCACAG GTG CTG CAG TTG GCT GTG CAG GCA CTG GTG CTA TAC TGT GGG CAC CAG CAG CTC
CGC GAA GGG ACC CTC ACT +1641
V L Q L A V Q A L V L Y C G H Q Q L

R E G D L T

FIGURE 8 - SUITE 2

+1642 GCC GGC AGC CTC GTC GCC TTC ATC CTC TAC CAG ACT AAA GCT GGC AGC TGC GTG CAG
GTGAGGTCAGGAGTGCCTCTGCCACCG +1729
A G S L V A F I L Y Q T K A G S C V Q

+1730
GATCCCCATGACTGTGGCCACATCCCCGTGTCCCCACCCTGGTGCTGTGGCTGGGGGTACATCCCCCATGTCCCTATCTGGGTGCTGTG
CATGCAG GCA CTG +1834

A L

+1835 GCG TAC TCC TAT GGT GAC CTT CTG AGC AAT GCA GTG GCC TGC AAG GTC TTT GAT TAC
CTG GAC TGG GAG CGA CCT GTG +1915
A Y S Y G D L L S N A V A A C K V F D Y
L D W E R P V

+1916 GGT GCT GGT GGC ACC TAT GTG CCC ACC AGA CTG CGG GGC CAC ATC ACC TTC CAT CGG GTG
TCC TTC GCC TAT CCC ACT CGC +1996
G A G G T Y V P T R L R G H I T F H R V
S F A Y P T R

+1997 CCT GAG CGC CTC GTC CTG CAA GAT GTC ACC TTC GAG CTG CGC CCC AGT GAG GTG ACG GCG
TTG GCG GGG CTG AAT GGC AGC +2077
P E R L V L Q D V T F G L R P S E V T A
L A G L N G S

+2078 GGG AAG AGC ACC TGC GTG GCA CTG CTG GAG AGA TTC TAT GAA CCT GGG GCC GAA GTG
CTG CTG GAC GGG GTG CCG CTG +2158
G K S T C V A L L E R F Y G P G A G E V
L L D G V P L

FIGURE 8 - SUITE 3

Figure 8

+2159 CGG GAC TAC GAG CAT CGC TAC CTG CAC CGC CAG
GTGAGGGGTGGGGGAAATGTAGCTGCACTGACACANTGCTGGGGCTGAACCTCTGCCTGG +2254
R D Y E H R Y L H R Q

+2255 GGGCAG GTG GCA CTG GTG GGG CAG GAA CCC GTG CTC TTC TCT GGC TCC ATT CGG GAT AAC
ATT GCC TAC GGG ATG GAG GAC +2335
V A L V G Q E P V L F S G S I R D N
A Y G M E D

+2336 TGC GAA GAG GAG ATC ATA GCA GCT GCA AGG GCT GCG GGT GCT TTG GGC TTC ATC TCT
GCA CTG GAG CAA GGC TTT GGC +2416
C E E E I A A A R A A G A L G F I S
A L E Q G F G

+2417 ACT G GTGAGTGTGGGGGAGCAAGGGGGGACCCGGGTGTCTGACCCCACTCATCCCCACCTCATCCTGCAG AC
GTA GGG GAG AGA GGG GGG CAG +2511
T
G E R G G Q

+2512 CTG TCA GCG GGG CAG AAG CAG CGC ATC GCC ATC GCC CGC GCT TTG GTG CGG CGT CCC ACC
ATC CTT ATC CTC GAC GAA GCC +2592
L S A G Q K Q R I A I A R A L V R R P T
I L I L D E A

+2593 ACC AGT GCT CTG GAT GGG GAC AGC GAT GCA ATG
GTGAGCACTGAGCAGTGGGTGGGGGGAGGGTCTG?CCCTGCAGTGCATGCTGATGGGCAGCTG +2608
T S A L D G D S D A M

+2609 TGTGTCCCTACAG CTA CAG CAG TGG GTG AGG AAC GGA GGG GAC CGG ACG GTG TTG TTT ATC ACC
CAC CAA CCA CGG ATG CTG +2769
L Q Q W V R N G G D R T V L F I T
H Q P R M L

```

+2770 GAG AAG GCA GAC CGC ATT GTG GTG CTG GAG CAT GGC ACG GTG GCT GAG ATG GGG ACA CCC
GCC GAG CTG AGG ACC CGC GGC +2850
      E K A D R I V V L E H G T V A E M G T P
A E L R T R G

+2851 GGA CCC TAC AGC CGG CTG TTA CAG CAC TGA
GAACCATGGAGCAGCTGGAGTGGCATGGGATGGGATATGGGGAGCAGTGACTGCCTTTGCTTCCAGC +2947
      G P Y S R L L Q H *

+2948 TGCAGGATGGGATGTTTTGGGATTTGTGTGGAATAAAGTGGAGATGCTTTGT
+2999      3'UT

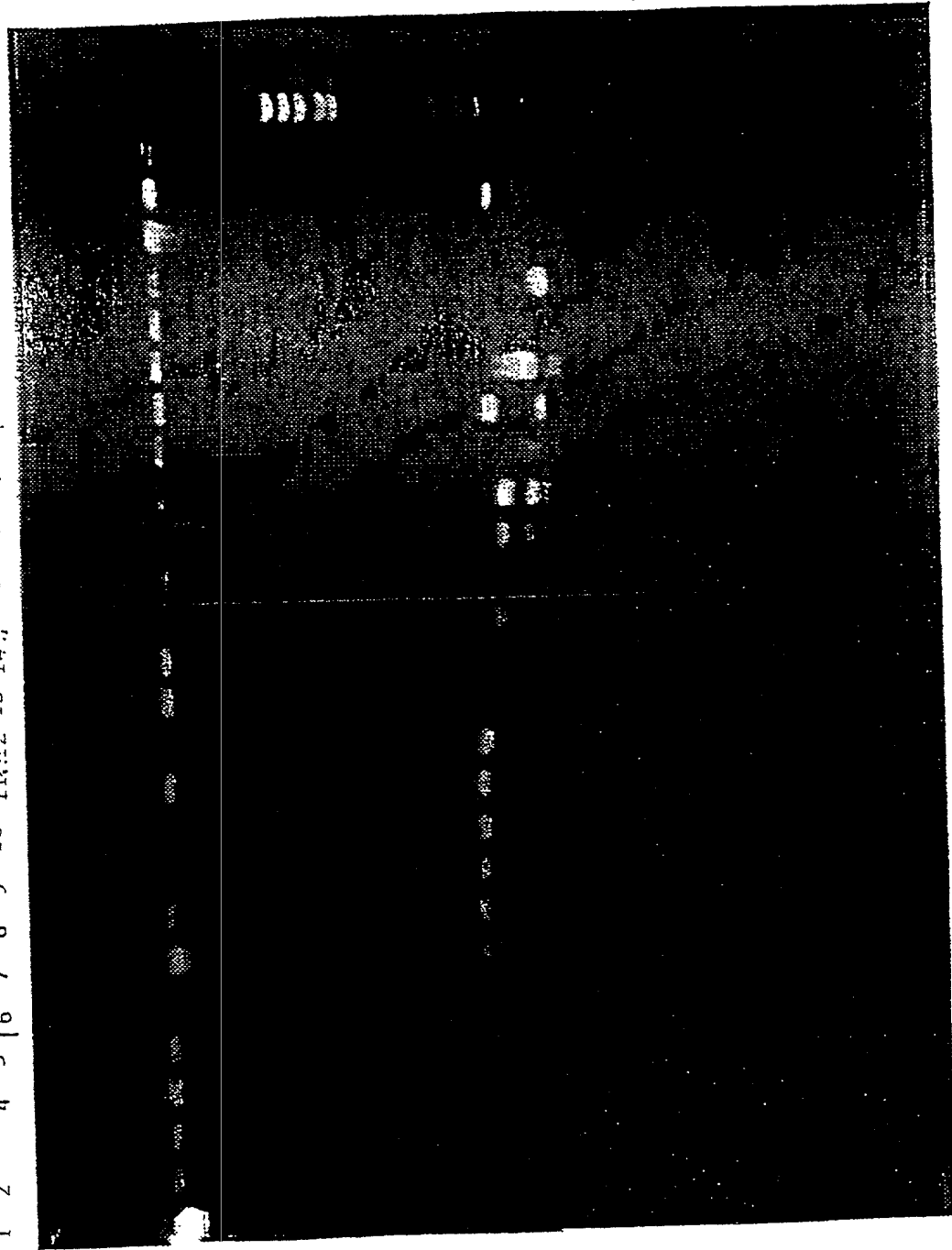
INTRON 2-3 : EF23(1B+1R)3R
INTRON 3-4 : EF23 2(1)
INTRON 4-5 : EF23 352H CON
INTRON 5-6 : EF23224RS
INTRON 6-7 : EF23(5B+5R)1R
INTRON 8-9 : EF23277B CON
INTRON 9-10: EF23 43RSR
INTRON 10-11: EF23 43RSR
INTRONS 1-2 ET 7-8 INEXISTANTS CHEZ LE POUET

```

FIGURE 8 - SUITE 5

FIGURE 9

B4
 1 2
 B5 B7
 4 5
 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 23 24 25 27
 B12 B13 B14 B15 BX
 B21



FEUILLE DE REMPLACEMENT (REGLE 26)

GTCCCTATTCCCATTGTGTCTCCTCACATCTGCCATCTCTTCTGTCCCCAT
 CTATGCTTTGTGCCCCCATCCCTTACCCCATCCCCACGTGTCCCTGTGG
 TGCCACCTCCACACGTGTCCCCGTGTCCCCACAGCGGGGCGGTGGCGCAA
 TAACACTGTGATGTGGCGCTGTGCGGGACGGAGCGACGGCGCTGCCCA
 TCCGTGCCACGTGCCAGCAGAGGGGACAGCGGGTGACGACGGCCGGGGG
 TGCCGAGACGCCTTCTGTCAGTGTGTGAGGTGGCACAGAATCTGCGGCG
 GAAGGGACAGCGCGGGGGTTGGCACGGGGTGAGTGTGACGAGTGTCCCC
 AAAGCGGGGAGGGGTGACCTGGGGTGGTGGCGGTGGGGTGTGGGGGAGTT
 GTAGAAATGGGGACCCCATTTGGTGTGGGGAGGTTTGGATAAGGGGTCCCC
 ATGGGTGGTGGCACATGGGGACATCCCATAGCCTGGGATCCCATGGTTGG
 GGCCATCCCGTACCTGGGATCCCCACATGGGAGGATGTCCCCCGCTGTCC
 CCATGGCAGTGTGAGGACACAGCTGGCAGAGCAGCTGTTGGATGATGAT
 GAGGACGTCCCCACGAGGAGCTTCTTCCCTGAGAGCTGGCTGTGGCGACG
 CATCCATGTTGCTGGCACTGCACGGTGTGTCCCCGTGTGTCCCCATGTCC
 CCATGTCCCCATGACTTTGTGTCCCCGTGTCCCCATCTCCCCATCTCCCC
 AGGCTCTCAGTGTCTCCCTGACTCCATCACTACGTGGGAGATTGAGGC
 AGTCGCCATCGTCCCTGGACATGGTGAGTGTACCCCCCTCCAATGGCCCT
 GCAGTGTCCCCCTGACATCCCCCTCGTGGTGTCCCCATGTCCCCCACGTC
 CCCAAGTTCTATGGTGTCCCCATGTCCCCCTCTCCCCCTCCCCCGGA
 ATGTCCCTGTGTCCCCGTGGTGTCCCTGCACTGCCCCGCAGTGATGAGGT
 CCTGGCAGGGGTGTGCGTGGCGGAGCCGACGGGTGACGGTGACACAGG
 ACGTGCGTGTGGCGCTTTGGCTGCCCCCAGCATCCGGCCCCCTAGAGCAG
 ATGCAGCTGCAGCCCCCTCATCCACAGCAGACTGCCCCGCAGCATCAACGT
 AAGCCCTATAGAGACCCCATAGGCACCCAGAGATACCTCTTCCCTCTA
 ATAAATACCACCTTTGCTTCCAATAGATAACCCTCCTGCCCCATAGGTACC
 CCTGTGCTCCATACTTGGCCTGCCACAGCATAACATACCCCTTCCCTCCA
 ACAGATATGCGTTGCCCCATAGATACTTCTTTCTGCCCTATAGATAACC
 CCTCATGCCCCACAGATTCCCGTTTCTTTCAATTGGTACCCCTGCCCC
 TCATATATCCCCCTCTACCCACAGGATACCCCTTAGACACCCGGTACCA
 CTTCTGCCCCATGGATACCCCTGTGGCACATAGATAACCGCTTCTGCCCC
 ACAGATACCCCTTCTACTCCACTGTCCCACAGCCCCCACTGCCCCATG
 GCCACCCATAGCCTGGTGGCATCGGGTGACAGTGACGGTGATGCAGGTGA
 CGGTGACACTGTGCGGAGTGGAGGGGTGTGCGCGGCGCTGGATGGGGTC
 CCCCAGATGCTGGAGCTGCCCCCGGGGAGGGCAGTGGCTGCACCCCTCAC
 TCTGGTGGCCCTCCACCCTGGGGACATCCCCATCACCATCACCGCCCGCG
 GGCCATGGGGGCTGGGGGACCGTGTACCCGAGTCTGCATGTGAGGTG
 AGATCAGTGGGGTCCCCTCCAGTCACCTGGGTACCTCTGGGGTCCCTTA
 AAGCCCTGCGACCTCCTGGACATTGTTGTCTTGTGAGCCTGCGGTACC
 CTGAATACTGGGGCTGTCACTTTGAGGTTTCATGGACACCATGTCCCTGTG
 TCCATGGTGGCCCTGGACATGTTGGTCTTATGGGATCTGGGGACATGGG
 GTCCTTGGTGGTCTGGATACTGCAGTTGTCTTTGTGGACACTATGTC
 CCCATGTCTTGGTGGGAATGGTGTTCATCCATTCCCGCAGCCTGAGGGAG
 AGCTGCACCTGGAGGAGACCTACATCCTGGACGAGATGGTGGGTGT
 GAGGACTGGGGGACACTGGGGAACTGGGGACGTGGGGCCGGACCCCTGTG
 GTGTGGTGTCCCTACAGATAAGCGGAGCCGGAGCCTGAAGCTGCCGGGGG
 ACGTCCCTGCAGAGATCGTCCCTGATGGGGACTTCAGCATGAGCATCCGT
 GTCAGTGGTGTGTGGGGATGGGGACATGGGGTGGGGACATGGGGGTGGGT
 ACTGGGAACGTGGTGGGGATGTGGTGGTGGGCATAGGGGACATGGGGACA
 TGGGAGGACATTTGTGGGGACATTGATGTCCATCCCTGATCATCTCTCT
 GTCCCTATGTCCCCATACCCATGTGTGTGGCCATGTCCGCACGCTGTGCC
 CCTGTGTGTGTCCCCGTGGTGTCCCCACATGTGCTCACATCCTTATTACA
 TCCCCACATCTCCTGTGTACAACCCCGTGTGCCCTGATGTGTGCCCTCC
 ACACATCCCCATGGGTGTCCCAATGTTCCCATGTCCCTCTGCTCATCCCC
 ATCCACATCCCCATGCTTGGGCTATCCCGAGCTTGGGGCAGTTGGG
 TCCTATCCCCATGTCCCCATGTCCCATATCTCCATACCCCTGTGACCCCA

FIGURE 10

10043450 011402

TATCCCTGTCCTTCAACTCCCCTCCCATCCCCACACCATCCCCATGTCTT
 CTGTCCCCACACCATCCCCATATCCCCCTGTCCCCCTGTCCCTGTCCCA
 GGCCGGGTGCCGGGCTGGGCACTGCAGGGCGCTCTGGGGATAGGGGACTC
 TCTGCTCCGCTCCCCCGGGGCTGTGGGGAGCAGTCCCTGATGTCAATGG
 CACCCACTGCTGCTGCTCTGCGCTTCCTGGATGAGAGCGAAGGGTGGGGG
 CAGCTGCCCCCAGGGCACCAGACAGCGCGGCCTCAGAACCCTGCAGCAGGG
 TGAGCTATGGGGCAGGTTGTGCTTTATGGGGTGGGCAATGCTTTATGGGG
 TGTGCAGTGCTCCAAGGGATGTGCAGTGCTTCATGGGGGATGCAGTGGGG
 TTTGATTTGATTTGATTTATGGGTTTGCATTTCTCCTCCGAGGATTGCAT
 CTCTCTATGGTGTGTTGCAATGGGATGTGCAGTGCTCCAGGTGGAGGTGCA
 GAGCCCTATGGGGGTGCAGTGCTGTGTAGGGGATGTCTGTGGTGTCCCCA
 ATGGTCTCTGATGTCCCCACAGGCTTCGAACGGGTGCAGAGCTTCGCGAA
 AAGTGACGGCTCCTATGGGGCATGGCTGCACCGGGACAGCAGCACCTGGT
 GAGGGGAGCGGGGATGATGTGGGACATGGGGATAGTGAGGGGATGTGGG
 GATGCTGGGGTATGGGGATGTGAGGACATCATAGGGACATGAGCGGTGGG
 GCCATGTGGATTTGGGGACGTGGTGACACGGTGTCTGGTGCAGGCTGAC
 GGCAGTGGTGTGCGTGTGCTGGCCCTGTCCCGGCCCTATTTGCCAGTGG
 CTGQCAGCGGCCCCGCTGCGTCCCTGCGGTGGGTGCTGGGGCAGCAGCGC
 CCAGATGGCGCCTTCTTGAGCACAGGGCTGTGGTGCACCGTGAGATGCA
 GGTGGGTGACACATCACTGCTGTGTGCAATGTCCCCATGCAGGATCTCCC
 CCTGCAATGTCCCCTGAAGGTCCCTGCAGGCTGACCCACATTACACTGT
 GTCACCTCACGTGTCCCCGTGTCCCCAGGGTGGTGTGGCAGACCCCGGCC
 GGAGGCCACCGTGTGCGTGACGGCCTTCGTGGTGGTGGCCCTCCATGGTG
 CCCGCGCTCTGCTGCCCCCGGACAGCCCTGAGCTGCCCCCTCCTGGTGAGT
 CCCATGTCCCCACCCCTGTGTCTTGGTCCCTCATATCCATGTGTCCCTGT
 GCCCCATCCCCCAAATCCCCACATCCCCCATATGTTCCCATACCCTGCTG
 TGTCCCCCGAGTGTCCCCCGTCTTTCATTCTCCACTATCCCCCGTATTC
 CCATATGTCCCCCTGTCCACAGTGTCCCCCTCATCCCTCTGTGTCCCCCT
 GTCCCCCAGTGTCCCCCAGTCCCTGTATGTCCCCATGTCTCCTAGTGTC
 CCCCATGTCCGTGTCTCCAGTATCCCCCATGCCTCCCCGTGTCTCTTCA
 TGCCCCACACTCCACGTCCCCACACTCCATGTCCCACTGCCACAGGACAA
 ATCCCTGTCCCGGGCCTCCACGTTCCTCCGGGGCCGCGTGGAGCAGTTGG
 GGACCTATGGGACAGCCATTACATCCTATGCATTGGCACTGGTGGACACC
 GCTCCTCCGGGGCCGCTATCCGGCGGTGGAACGTCTGCGGGGCATGGCCCG
 GAGCGCCACGGTGCGTCTGTCTGTCCCCATGGGGTGGTGGCACCTCT
 GTCCCCATGGCTGCCTCCTGGACCCCTCTGTCCCCTCCTTCAGATTCACT
 CTCATTGCAATCCTTCAATTTTATTCTCCCTCAAACCTCTTCTTTGTA
 TTCTTCACATTCACTCCTATTCAAATTGCTCTCCTTCTGTCTGTCTTC
 TTCAAATTCTTCTTCAATTTTGTCTCCTGATTAATTCTCTTAAATTA
 CTCTCGATCAAGTTCTGCAGATTCTGTTCCACTTCGGATGGATTCTTCTCC
 AAACCTGTTCTTCAGATTCACTCTCCTTCAATTTCTGTTCTTGTAAATTA
 CTTCTTCAGAGTGATTCTTCAAACCTCTTCTTCATGTTCTCTTCAAGTCCA
 TTCCCTGCACTGACTCCGGGTGCTCAGGACCCCCCGTGACCCCATATGA
 CCCCATATGAACCCCCCATGACCTCCACAAAACCATATGACCCCGTGACC
 TCCCATGACCCCTCATGACCCCATATGACCCCATGACCCCATCCCTGT
 GCAGGTGGCCGTGCAACCTTCTGGCCATCCGGTGGCCCCCGCAGCCACGGT
 GGAGGCGACGGGTACGCCCTTCTGGCACTGCTGCAGAGCCGCGACATCG
 CCGGGGCTGCGAGGGCGGCACGGTGGCTCCGACAGCAGAGCAATTACGGG
 GGTGGCTTCCACTCCACGCAGGTGGGTGGGGGTCACTGACCCCCCGGGTG
 CCTCGGGGTGGGGGTGATTTGATCCCCAGGTACCTCTTTGGTGGCTGTGT
 CCCCCAACCTGCTTGGTGTCCCCCAGGACACGCTGGTGGCCCTGGAGGCG
 CTGGCCCAGATGTGGCTGCACTGGGGCCGTGGGAACACAATGGGGCTGAA
 CCTGGGGCTCTCCTGGCCGGGGGGTGGCCGGGGGAGGGCTGGTGGCACTC
 AGGTTATGCTGAAGCCGGGGCTGGACCCCTGGAGCAGGAGCTGGAGGTG
 GGGACATGGCGGGATGTGGGGACAGGAGGATGTGAGGACACTGGGGACA

FIGURE 10

SUITE 1

FEUILLE DE REMPLACEMENT (REGLE 26)

TGTCTGGACTTGGTAGGATGTAACATGAAGACACTGGGGACATGGTAGGA
 CATGGGGGACATGAGAACACGGGATGTGGGGGACATGGTAGGACATGATG
 GACACAGGGCTTTGGGGTCTTGGGTCTCGCTCTGTCCCATGTCCCCA
 GGTGCCTCTGGGCAGCCCAGTGACAGTGCAGGTGGAGGGACACGGCGAAG
 GGACGCTGACGGTGGGTGGCTGCATGGACATTGGTGTCTCTCAAGACC
 GATGTCCCCCTCACAACTCCCCCTCATGGTGTCCCCCTCATGCTGCCACGGT
 GTCCCCCTGCTGTCCCCTCATGGTGTACGCTGTCCCCAGGTGCTCCGCCA
 GTTCCGCCTGCTGTACCTCCGAACGCCACGTGCCAGGCGCTGCACCTGG
 AGGTGGCCATCACCGGCCCCCATCCTGTACCATGGTGAGGCCCCACCCAAA
 GGCCCCGCCCCCTTTTCTCGCGGGGGGCGTGCCCTCAACCCTGTTTTGC
 ATATCCCAACCCCCAGCAGATGAGGACTACGAGGACTACGAGGACTACGA
 GGAGGCGGAGCCTAAGGAGGGGGAGGAGCCTACGGAAGGGGCAGTGCCCG
 TGGAAGGGGCGGGGCCAGCAGATGACCCCGCCCCCTCAGCCCCGTGTCC
 TTATGGGATGCCCGTAAGCGGCAACGCCGCAGCACACATAACCCTGCCCA
 CGAGGTGGCCTTCTGGTCTGCTTCCGGTGAGGGGCGGAACCTTCTGTCC
 CTGGGGGCGGGTCTTCTGCTGATGGGCGTGGCTTATTGCTGAGGGGCGT
 GGCTGTGTAGGCGGAGCCAGGGGTGGCACTGACTGGGATGGCGGTGG
 TGGAGATCACTCTGCTCAGTGGCTTCTCACCCCATAGAGCTGACCTGGAC
 AAGGTAGGGGCCCAGGGGGACTTGTGGGACATGTTGGGGGGTTGAGGGGA
 GTTATGGGGTGTGGGGTTTGGGGGTGTTGGAGTTGTTGAGGTGGCAGAAT
 GTTTGGGTGGAGTCATGGGATATGGGGCTATTGGGGTTTGGAGGTGTTG
 TGATGTTGGGAAACATTGAATTGGGGTTGTTGAGTTTGGGGTGTGGGG
 TGTGCGGGTGCAGAGCTGCAGCTGCTGGGTGGAGTATTAAGGTGTTGGG
 ATGTTGGGGTGTGGATGGCTTGGATGCGGGTGTGGGGTGGGCACGTAT
 CTGGGTGCTGCTGTCCACACAGCTGCGGGACGTGGTGGATCACTGGAT
 CAGTCACTATGAGTTGGAAGGAAACCAGTTGGTGTATACCTGGATGAGG
 TGTGTCTTCCCGTGTCAACCCTATAACCCAGTGGCCCCATGTTCTCATAT
 CCCCCATGTCCCCGTGTCCCCACACCATATCCCATTTCTCCCCACACATCC
 CCGTGTTCACACAGTGTCTCTATTCTGTCCCTGTCCCCAGGTCCCCCC
 CGAGCGGCAGTGTCTCAGTTTTGGGGCCACCCAGGACGCGGCTGTGGGTC
 ACATGCAGCCGGCAATGGCAGCCATCTATGACTACTATGAGCCTGGTGGG
 TGGGGCCTTCAGTGGGAGGGGCTAAATGGGTGGTGGTCTTCATGGGTGT
 GACCATTGGAGGAGGCGTGGCCGATCTGACCCCTCCATGCCCCATCCAGG
 ACAGCGCTGCACCGTCTTCTACAACGCCCCCCCAAAGGAGCAGCACCATCG
 CCACACTGTGCTCCCCCAAAATCTGTGAATGCGCCCCAAGGTAGGACCCCA
 CTGTGACTCCATATGTAGGGCCCCCATCCAGTGAACCCCCACATCCTCCT
 CCTAATTTTTGAAGATCTGGGGGTGAAATTATGGGGTTTATAGGGGAGCG
 TGGTTGAGTGACATGCAGGACATGGAGGGAACCCACACCAAGAACCTTGT
 GTTTTGGGTCCCTGATGATGTTGGGAGATCCTATTGATGTTGGTGGTCCC
 CAGGGGGGTGTCCCCAAGCCCAAAGGAGGACACAGGAGGTGACAGCTGAT
 GACCGCCATGACTTTGCCTGCTACAGCCCCCGCTGGACTATGGTGAGAT
 CCCAAATCACTGCACCTCAAACCTGACCCCAAATTTGGCTGCATCCCGAAC
 CCCAACTGCCCTAAATCCCATCTGCTGCCCCCTGAGTCCCACAGCTGCACA
 CTGTACCCACAAAGTGGCCCCCTGAAGCCTAAAAACATTACAGAGGATTTT
 GTAGTTTTCTCCCTGTACCCAGTTGTCCCTCTGACCCCAAGAACCCAC
 AGCTGCCCTATGCTGTCCCTGCCCCGCCATAACTCCTCTGATACAATAAC
 CCCCCTGACCCCATCTTTATGACCTCCATGACCTTTGACCCCCAGCACTG
 GTGGTTGGGGTGTGTCCAGAGTGAGATAGGGGCTTTTGTGGCGTTTGA
 GACGGAATCAAGGAGGTGCTGCTTGAAGGTGAGACTGAGGGTAGTGGGA
 CGGACTGGAAGGTGAGAATGGGAGCACTGGGAGAGGCAGGGAGTACTGAG
 AGGGACTGGGAATGACTGGAAATTGAGACTGGGTGGACTGGGAACCTCTGG
 TAGAGACTGAATGGGTATACTGGGAACACTGGAAGAAGTTGTGGGATGAG
 AAGAGGATGCTGGGATAGGAGACCCCCCCCCCTTGTGCTAGGGGGGTCTCT
 CAGCCATACTGGGCACAATATGAGAGTACTGGGTGGTACTGGGAAAGCT
 GGGAGGACTCATACTGGTCTCTACTGCTGAGGCGGAGGAGGAGGAGGAGG

FEUILLE DE REMPLACEMENT (REGLE 26)

FIGURE 10

SUIITE 2

10043160 011402

CCCCCTGGGGAGCGGAGGCGGCTGCTGGTGCGGAAGAGCTGCCCACTGCGC
CTGCAACTCCACAACATCTACCTGGTGATGGGGGGCAGCGGGAGGACGCG
GGACCCTGAGGGGCGGTGAGAAGGGGCTGTGCCCCATGTCCACATGTCCC
TGTGTTCTCATGTTCCCATGTCCCATATCCCAGTGTTCCCTAACCCCATAT
CCTTGACCTTGAGCCCATACCCCTGATATCCCTGACCCTGTCCCCATTCTC
AGCCCCCAGTTCCTGCTGGGCCCCCACTCATGGTTGGAGGAGGTGCCATC
CCCTGGACGCTGTAAGGCCACAAGGTTGCGGGGTTACTGCGCCCAACTGC
AGGAGTTCGCGACCCGCTGAGCCAACTGGGCTGCCAGCTGTGAGCCCCT
GGGAGCCACTGGGAGCATGTTGGGTGCAGCTGGGACCATTCTGGGGGTGA
AGTGGTACCACTGTTGGATCAGTTGGGATCAATTGGGAATAAACTAGTGT
TGACTGGGACCGTGTTGTGACCAACTGGAAGTGTGTTGGAAGAACTGAG
AGCTGCTGGGGTTGAGTGGGAGCAACTGGAAGTGTGTTGGAACAAACAGG
GGACCAACTGGGATCACACTGTGGTCAGCTGGGATCACACTGGGTCAAAA
AAGATCACAGTGGCCCAATTGGGGTCATACTGGGGTGAGCTGGGATCAGA
ACGAGTTTAAATAAACGTACAGTCGTCCGAGCCACCACAGAGTCAGCCCTC
CAGCGGCGCAGAGCGGCGCAGCGCGCACTGGCTGCCCGCGGTAAAGCGGAT
GTGACGTCACTTCGCGGCGCGCTATTGAACTCCAGCAGCGCCCCGCGGA
GCGCCCCAATGCCGCGGCCCAAACCGCGCAGCCCCCGCGCGGGGCGCG
CCCCCCCCGCGCGCCCCCCCCCGCCACCCCCCGCGCGGCTCGCGGTGAGTG
CAGCCCGTAGGAGTGCGGAGTGTGGGGGCGGGGGGGGGGCGTCTGGAGC
GGAGCCTTTATCACCGCTGTTTTCCCGATTTCCCGTCTTTTCGCCCCGT
TTCAGCCCGCCGGTACCGGCCCGGTGAGAGGCGCTGCGGGAGATCCGCC
GCTATCAGAGCAGCACCGCTCTGCTGCTGCGCCGCCAGCCCTTCGCGCGC
GTGGTAACGGGACTGCCCCGGAACGGGACACCCCCCAACCCCCCAACGG
GACCATCCCCCACGGATGGATCCCCCCCCACACACATCCAACGTGGGAC
CCCCCGCCCCAAAATGAGATCTCAACGTGAGATCTGGGGGCCTCAAAATG
AGACACTCTCCCCCTCCCCCAACGGAACACCCCCGAAAATGGGACCACAC
ATAAAAGTGGGACTCCCCCTCTCCCCCGCGCCCGTCAAAATGGAACAC
CCCCAACTGGACCTTTCAAAAAATAACATTCCCCCTCCCCCAAAAATGGG
ACTTACCACAAAGTGGGATCTTCCCCCAAAATGAACACCCCTCAAAATG
AGACCCCTCGGACCCCCCCCCAACCCCTCTGCACCCATCNGCCGTCTGTGA
CGGAAGGGAAAGGCTGTAGGGTACATCTACCCTTATTTCTTGGGTTTGTG
TTTTGTTTTGTTGTTATTTAGAAGCAAAACCAAGACAACAAAGCCCAGCC
AATGCCATTTCTGGCAGTGGACGCAGGCGCAGGCGGGTTGGTCACAAAG
CAAGAAGTTGCTGCGGGACTTTGTGTTTTGGGGCCGTTCTCGTGAACCT
CTGAGCCATGGATGAGGAAATTACTTATGCTGATTTAAGGCATCCTACGG
GCAGTTTGCCTCCTGCTAAGCGGCAGCGCGGTAAAGGATGCTCTGTGTGG
TGGGTGCTCACCGCAGGCTTGGTTTGGGGGCTTGCTGTTCTCTGAGAAAC
ACCAGCAATGCTGGTTGGGTTCTGGGTCCACCCTGGCTTGTATGGGGGAG
TAAAGGAAGGGGTGGGGGAGAAGGAAGCCTGGGAATGGCCAGAGGTGTGG
TGGTTTTT

FIGURE 10

SUITE 3

10043450.01402

GTCCCTATTCCCATTTGTGTCCTCACATCTGCCATCTCTTCTGTCCCCAT
 CTATGCTTTGTGCCCCCATCCCTTACCCCATCCCCACGTGTCCCTGTGG
 TGCCACCTCCACACGTGTCCCCGTGTCCCCACAGCGGGGCGGTGGCGCAA
 TAACACTGTGATGTGGCGCTGCTGCCGGGACGGAGCGACGGCGCTGCCCA
 TCCGTGCCACGTGCCAGCAGAGGGGACAGCGGGTGACGACGGCCGGGGGC
 TGCCGAGACGCCTTCCTGCAGTGCTGTGAGGTGGCACAGAATCTGCGGGC
 GAAGGGACAGCGCGGGGGGTGGCACGGGGTGAGTGTGAGCAGTGTCCCC
 AAAGCGGGGAGGGGTGACCTGGGGTGGTGGCGGTGGGGTGTGGGGGAGTT
 GTAGAAATGGGGACCCCATTTGGTGTGGGGAGGTTTGGATAAGGGGTCCCC
 ATGGGTGGTGGCACATGGGGACATCCCATAGCCTGGGATCCCATGGTTGG
 GGCCATCCCGTACCTGGGATCCCCACATGGGAGGATGTCCCCCGCTGTCC
 CCATGGCAGTGATGGAGGCACAGCTGGCAGAGCAGCTGTTGGATGATGAT
 GAGGACGTCCCCACGAGGAGCTTCTTCCCTGAGAGCTGGCTGTGGCGACG
 CATCCATGTTGCTGGCACTGCACGGTGTGTCCCCGTGTGTCCCCATGTCC
 CCATGTCCCCATGACTTTGTGTCCCCGTGTCCCCATCTCCCCATCTCCCC
 AGGCTCTCAGTGCTGCTCCCTGACTCCATCACTACGTGGGAGATTGAGGC
 AGTCGCCATCGTCCCTGGACATGGTGAGTGTACCCCCCTCCAATGGCCCT
 GAGTGTCCCCCTGACATCCCCCTCGTGGTGTCCCCATGTCCCCACGTC
 CCAAGTTCTCTATGGTGTCCCCATGTCCCCCTCTCCCCCTCCCCCGGA
 ATGTCCCTGTGTCCCCGTGGTGTCCCTGCACTGCCCCGCAGTGATGAGGT
 CCTGGCAGGGCTGTGCGTGGCGGAGCCGCAGCGGGTGACGGTGACACAGG
 ACGTGCGTGTGGCGCTTTGGCTGCCCCCAGCATCCGGCCCTAGAGCAG
 ATGCAGCTGCAGCCCCCTCATCCACAGCAGACTGCCCCGCAGCATCAACGT
 AAGCCCTATAGAGACCCCATAGGCACCCAGAGATACCTCTTTCCCTCTA
 ATAAATACCACCTTTGCTTCCAATAGATAACCCTCCTGCCCCATAGGTACC
 CCTGTGCTCCATACTTGCCCTGCCACAGCATACATAACCCCTTTCCCTCCA
 ACAGATATGCGTTGCCCCATAGATAACCTTCTTTCTGCCCTATAGATAACC
 CCTCATGCCCCACAGATTCCCGTTTCCTTTCAATTGGTACCCCTGCCCC
 TCATATATCCCCCTCTACCCACGGATACCCCTTAGACACCCGGTACCA
 CTTCTGCCCCATGATAACCCCTGTGGCACATAGATAACCGCTTCTGCCCC
 ACAGATAACCCCTTCTACTCTGCTGCCCCACAGCCCCCACTGCCCCATG
 GCCACCCATAGCCTGGTGGCATCGGGTGACAGTGACGGTGATGCAGGTGA
 CGGTGACACTGTGCGCAGTGAGAGGGGGTGTGCGCGCGCTGGATGGGGTC
 CCCCAGATGCTGGAGCTGCCCCCGGGGAGGGCAGTGGCTGCACCCCTCAC
 TCTGGTGGCCCTCCACCCTGGGGACATCCCCATCACCATCACCGCCCGG
 GGCCATGGGGGCTGGGGGACCGTGTCAACCGAGTCCTGCATGTGAGGTG
 AGATCAGTGGGGTCCCCTCCAGTCACCTGGGTACCTCTGGGGTCCCTTA
 AAGCCCTGCGACCTCCTGGACATTGTTGTCCTTGTGAGCCTGCGGTCAAC
 CTGAATACTGGGGCTGTCACTTTGAGGTTGATGGACACCATGTCCCTGTG
 TCCATGGTGGCCCTGGACATGTTGGTCTTATGGGATCTGGGGACATGGG
 GTCCTTGGTGGTCTGGATACTGCAGTTGTCTTTTGTGGACACTATGTC
 CCCATGTCTTGGTGGGAATGGTGTATCCATTCCCGCAGCCTGAGGGAG
 AGCTGCACCTGGAGGAGAGCACCTACATCCTGGACGCAGATGGTGGGTGT
 GAGGACTGGGGGACACTGGGGAACTGGGGACGTGGGGCCGGACCCCTGTG
 GTGTGGTGTCCCTACAGATAAGCGGAGCCGGAGCCTGAAGCTGCCGGGGG
 ACGTCCCTGCAGAGATCGTCCCTGATGGGGACTTCAGCATGAGCATCCGT
 GTCAGTGGTGTGTGGGGATGGGGACATGGGGTGGGGACATGGGGGTGGGT
 ACTGGGAACGTGGTGGGGATGTGGTGGTGGGCATAGGGGACATGGGGACA
 TGGGAGGACATTTGTTGGGGACATTGATGTCCATCCCTGATCATCTCTCT
 GTCCCTATGTCCCCATACCCATGTGTGTGGCCATGTCCGCACGCTGTGCC
 CCTGTGTGTGTCCCCGTGGGTGTCCCCACATGTGCTCACATCCTTATTACA
 TCCCCACATCTCCTGTGTACAACCCGTGTGCCCTGATGTGTGCCCCCTCC
 ACACATCCCCATGGGTGTCCCAATGTTCCCATGTCCCTCTGCTCATCCCC
 ATCCACATCCCCATGCCATGCCCTATCCCCATCCCCACGTTCCCCCGATTCCCA
 TCCTATCCCCATGTCCCCATGCCCTATCCCCATCCCCATCCCCATCCCCAT

FIGURE 10

SUITE 4

20043150 "01402

TATCCCTGTCCTTCAACTCCCCTCCCATCCCCACACCATCCCCATGTCTT
CTGTCCCCACACCATCCCCATATCCCCCTGTCCCCCTGTCCCTGTCCCA
GGCCGGGTGCCGGGCTGGGCACTGCAGGGCGCTCTGGGGATAGGGGACTC
TCTGCTCCGCTCCCCCGGGGCTGTGGGGAGCAGTCCCTGATGTCAATGG
CACCCACTGCTGCTGCTCTGCGCTTCCCTGGATGAGAGCGAAGGGTGGGGG
CAGCTGCCCCCAGGGCACCACAGCGCGGCCCTCAGAACCCTGCAGCAGGG
TGAGCTATGGGGCAGGTTGTGCTTTATGGGGTGGGCAATGCTTTATGGGG
TGTGCAGTGCTCCAAGGGATGTGCAGTGCTTCATGGGGGATGCAGTGGGG
TTTGATTTGATTTGATTTATGGGTTTGCATTTCTCCTCCGAGGATTGCAT
CTCTCTATGGTGTGTTGCAATGGGATGTGCAGTGCTCCAGGTGGAGGTGCA
GAGCCCTATGGGGGTGCAGTGCTGTGTAGGGGATGTCTGTGGTGTCCCA
ATGGTCTCTGATGTCCCCACAGGCTTCGAACGGGTGCAGAGCTTCCGCAA
AAGTGACGGCTCCTATGGGGCATGGCTGCACCGGGACAGCAGCACCTGGT
GAGGGGAGCGGGGATGATGTGGGGACATGGGGATAGTGAGGGGATGTGGG
GATGCTGGGGTATGGGGATGTGAGGACATCATAGGGACATGAGCGGTGGG
GCCATGTGGATTTGGGGACGTGGTGACACGGTGTCTGGTGCAGGCTGAC
GGCACTGGTGCTGCGTGTGCTGGCCCTGTCCCGGCCCTATTGGCAGTGG
CTGCCAGCGGCCCGCTGCGTCCCTGCGGTGGGTGCTGGGGCAGCAGCGC
CCAGATGGCGCCTTCTTGAGCACAGGGCTGTGGTGCACCGTGAGATGCA
GGTGGGTGACACATCACTGCTGTGTGCAATGTCCCCATGCAGGATCTCCC
CCTGCAATGTCCCTGAAGGTCCCTGCAGGCTGACCCACATTACACTGT
GTCACTCACGTGTCCCGTGTCCCCAGGGTGGTGTGGCAGACCCCGGCC
GGAGGCCACCGTGTGCTGACGGCCTTCGTGGTGGTGGCCCTCCATGGTG
CCCGCGCTCTGCTGCCCCCGGACAGCCCTGAGCTGCCCTCCTGGTGAGT
CCCATGTCCCCACCCCTGTGTCTTGGTCCCTCATATCCATGTGTCCCTTGT
CCCCCATCCCCAAATCCCCACATCCCCCATATGTTCCCATACCCCTGTG
TGTCCTCCCGAGTGTTCCTCCCGTCTTTCATTCTCCACTATCCCCCGTATTC
CCATATGTCCCCCTGTCCACCAGTGTCCCTCATCCCTCTGTGTCCCCCT
GTCCCCCAGTGTCCCCCAGTCCCTGTATGTCCCCATGTCTCCTAGTGTC
CCCCATGTCCGTGTCTCCAGTATCCCCCATGCCTCCCCGTGTCTCTTCA
TGCCCCACACTCCACGTCCCCACACTCCATGTCCCACTGCCACAGGACAA
ATCCCTGTCCCGGGCCTCCACGTTCTCCGGGGCCGCGTGGAGCAGTTGG
GGACCTATGGGACAGCCATTACATCCTATGCATTGGCACTGGTGGACACC
GCTCCTCCGGGGCCGCATCCGGCGGTGGAACGTCTGCGGGGCATGGCCCCG
GAGCGCCCACGGTGCGTCTGTCTGTCCCCATGGGGTGGTGGCACCTCT
GTCCCCATGGCTGCCTCCTGGACCCCTCTGTCCCCCTCCTTCAGATTCACT
CTCATTGCAATCCTTCAATTTATTCTCCCTCAAACCTCTTCTTCTTTGTA
TTCTTCACATTCATTCTTCAATTTCAATTTGCTCTCCTTCTGTCTGTTCTTC
TTCAAATTCTTCTTCAATTTTGTCTCCTGATTAATTCTCTTAAAATTAA
CTCTCGATCAAGTTCTGCAGATTCTGTTCCACTTCGGATGGATTCTTCTCC
AACTGTTCTTCAGATTCACTCTCCTTCAATTTGTTCTTGTAAATTAATT
CTTCTTCAGAGTGATTCTTCAAACCTCTTCTTCATGTTCTCTTCAAGTCCA
TTCCCTGCACTGACTCCGGGTGCTCAGGACCCCCCGTGACCCCATATGA
CCCCATATGAACCCCCCATGACCTCCACAAAACCATATGACCCCGTGACC
TCCCATGACCCCTCATGACCCCATATGACCCCCCATGACCCCATCCCTGT
GCAGGTGGCCGTGCAACCTTCTGGCCATCCGGTGGCCCCGCGACACGGT
GGAGGCGACGGGTACGCCCTTCTGGCACTGCTGCAGAGCCGCGACATCG
CCGGGGCTGCGAGGGCGGCACGGTGGCTCCGACAGCAGAGCAATTACGGG
GGTGGCTTCACTCCACGCAGGTGGGTGGGGTCACTGACCCCCCGGGTG
CCTCGGGGTGGGGGTGATTTGATCCCCAGGTACCTCTTGGTGGCTGTGT
CCCCAACCTGCTTGGTGTTCGCCGAGGACACGCTGGTGGCCCTGGAGGCG
CTGGCCCAGATGTGGCTGCACTGGGGCCGTGGGAACACAATGGGGCTGAA
CCTGGGGCTCTCCTGGCCGGGGGTGCCCGGGGAGGGCTGGTGGCACTC
AGGTTATGCTGAAGCCGGGGCTGGAGCCGCTGGAGCAGGAGCTGCAGGTG
GGGACATGGCGGGATCTGGGGACACAGGGATGTGAGGACACTGGGGACA

FEUILLE DE REMPLACEMENT (REGLE 26)

FIGURE 10

SUITE 5

TGTCTGGACTTGGTAGGATGTAACATGAAGACACTGGGGACATGGTAGGA
 CATGGGGGACATGAGAACACGGGATGTGGGGGACATGGTAGGACATGATG
 GACACAGGGCTTTGGGGTCTTGGGTCTCGCTCTGTCCCCATGTCCCCA
 GGTGCCTCTGGGCAGCCCACTGACAGTGCAGGTGGAGGGACACGGCGAAG
 GGACGCTGACGGTGGGTGGCTGCATGGACATTGGTGTCTCTCCAAGACC
 GATGTCCCCCTCACAACCTCCCCCTCATGGTGTCCCCCTCATGCTGCCACGGT
 GTCCCCCTGCTGTCCCATCATGGTGTGACGCTGTCCCCAGGTGCTCCGCCA
 GTTCCGCCTGCTGTGACCTCCGAACGCCACGTGCCAGGCGCTGCACCTGG
 AGGTGGCCATCACCGGCCCCATCCTGTACCATGGTGAGGCCCCACCCAAA
 GGCCCCGCCCCCTTTCTCGCGGGGGGCGTGCCCTCAACCCTGTTTTGC
 ATATCCCAACCCCCAGCAGATGAGGACTACGAGGACTACGAGGACTACGA
 GGAGGCGGAGCCTAAGGAGGGGGAGGAGCCTACGGAAGGGGCAGTGCCCG
 TGGAAGGGGCGGGGGCCAGCAGATGACCCCGCCCCCTCAGCCCCGTGTCC
 TTATGGGATGCCCCGTAAGCGGCAACGCCGCAGCACACATAACCCTGCCCA
 CGAGGTGGCCTTCTGGTCTGCTTCCGGTGAGGGGCGGAACCTTCTGTCC
 CTGGGGGCGGGTCTTCTGCTGATGGGCGTGGCTTATTGCTGAGGGGCGT
 GGCCTGTTGTAGGCGGAGCCAGGGGTGGCACTGACTGGGATGGCGGTGG
 TGGAGATCACTCTGCTCAGTGGCTTCTCACCCCATAGAGCTGACCTGGAC
 AAGGTAGGGGCCCAGGGGGACTTGTGGGACATGTTGGGGGGTGTAGGGGA
 GTTATGGGGTGTGGGGTTTGGGGGTGTTGGAGTTGTTGAGGTGGCAGAAT
 GTTTGGGTGGAGTCATGGGATATGGGGCTATTGGGGTTTGAGGGTGTG
 TGATGTTGGGAAACATTGAATTGGGGTTGTTGAGTTTGAGGGTGTGGGG
 TGTGCGGGTGCAGAGCTGCAGCTGCTGGGTGGAGTATTAAGGTGTTGGG
 ATGTTGGGGTGTGGATGGCTGGATGCGGGTGTGGGGTGGGCACGTAT
 CTGGGTGCTGCTGTCCCAACAGCTGCGGGACGTGGTGGATCACTGGAT
 CAGTCACTATGAGTTGGAAGGAAACAGTTGGTGTATACCTGGATGAGG
 TGTGTCTTCCCGTGTACCCCTATAACCCAGTGGCCCCATGTTCTCATAT
 CCCCCATGTCCCCGTGTCCCCACACCATATCCCATTTCTCCCCACACATCC
 CCGTGTTCACCACGTGTCTCATTCTGTCCCTGTCCCCAGGTCCCCC
 CGAGCGGCAGTGTCTCAGTTTTGGGGCCACCCAGGACGCGGCTGTGGGT
 ACATGCAGCCGGCAATGGCAGCCATCTATGACTACTATGAGCCTGGTGGG
 TGGGGCCTTCAGTGGGAGGGGCTAAATGGGTGGTGGTCTTCATGGGTGT
 GACCATTGGAGGAGGCGTGGCCGATCTGACCCCTCCATGCCCCATCCAGG
 ACAGCGCTGCACCGTCTTCTACAACGCCCCCAAGGAGCAGCACCATCG
 CCACACTGTGCTCCCCAAAATCTGTGAATGCGCCCCAAGGTAGGACCCCA
 CTGTGACTCCATATGTAGGGCCCCCATCCAGTGAACCCCCACATCCTCCT
 CCTAATTTTTGAAGATCTGGGGGTGAAATTATGGGGTTTATAGGGGAGCG
 TGGTTGAGTGACATGCAGGACATGGAGGGAACCCACACCAAGAACCTTGT
 GTTTTGGGTCCCTGATGATGTTGGGAGATCCTATTGATGTTGGTGGTCCC
 CAGGGGGGTGTCCCCAAGCCCCAAGGAGGACACAGGAGGTGACAGCTGAT
 GACCGCCATGACTTTGCCTGCTACAGCCCCCGCGTGGACTATGGTGAGAT
 CCCAAATCACTGCACCTCAAACCTGACCCCAAATTGGCTGCATCCCGAAC
 CCCAACTGCCCTAAATCCCATCTGCTGCCCTGAGTCCACAGCTGCACA
 CTGTACCCACAAAGTGGCCCCCTGAAGCCTAAAAACATTACAGAGGATTTT
 GTAGTTTTCTCCCTGTACCCCAAGTTGTCCCTCTGACCCCAAGAACCCAC
 AGCTGCCCTATGCTGTCCCCTGCCCGCCATACTCCTCTGATAACAATAAC
 CCCCCTGACCCCATCTTTATGACCTCCATGACCTTTGACCCCAAGCACTG
 GTGGTTCCGGTGTGTCCCAGAGTGAGATAGGGGCTTTTGTGGCGTTTGA
 GACGGAATCAAGGAGGTGCTGCTTGAAGGTGAGACTGAGGGTAGTGGGA
 CGGACTGGAAGGTGAGAATGGGAGCACTGGGAGAGGCAGGGAGTACTGAG
 AGGGACTGGGAATGACTGGAAATTGAGACTGGGTGGACTGGGAACCTCTGG
 TAGAGACTGAATGGGTATACTGGGAACACTGGAAGAAGTTGTGGGATGAG
 AAGAGGATGCTGGGATAGGAGACCCCCCCCCCTTGTGCTAGGGGGGTCTCT
 CAGCCATACTGGCACAATATGAGAGTATACTGGGTGGTACTGGGAAAGCT
 GGGAGGACTCATACTGGT

FIGURE 10

SUIITE 6

20443160-0340

REUILLE DE REMPLACEMENT (REGLE 26)

CCCCTGGGGAGCGGAGGCGGCTGCTGGTGCGGAAGAGCTGCCCACTGCGC
 CTGCAACTCCACAACATCTACCTGGTGATGGGGGGCAGCGGGAGGACGCG
 GGACCCTGAGGGGCGGTGAGAAGGGGCTGTGCCCCATGTCCACATGTCCC
 TGTGTTCTCATGTTCCCATGTCCCATATCCCAGTGTTCCCTAACCCCATAT
 CCTTGACCTTGAGCCCATACCCTGATATCCCTGACCCTGTCCCCATTCTC
 AGCCCCCAGTTCCTGCTGGGCCCCCACTCATGGTTGGAGGAGGTGCCATC
 CCCTGGACGCTGTAAGGCCACAAGGTTGCGGGGTTACTGCGCCCAACTGC
 AGGAGTTCCGCACCCGCTGAGCCAAGTGGGCTGCCAGCTGTGAGCCCCCT
 GGGAGCCACTGGGAGCATGTTGGGTGCAGCTGGGACCATTCTGGGGGTGA
 ACTGGTACCCTGTTGGATCAGTTGGGATCAATTGGGAATAAACTAGTGT
 TGAAGTGGGACCGTGTGTTGACCAACTGGAAGTGTGTTGGAAGAACTGAG
 AGCTGCTGGGGTTGAGTGGGAGCAACTGGAAGTGTGTTGGAACAAACAGG
 GGACCAACTGGGATCACACTGTGGTCAGCTGGGATCACACTGGGTCAAAA
 AAGATCACAGTGGCCCAATTGGGGTCATACTGGGGTGAGCTGGGATCAGA
 ACGAGTTTAAATAAACGTACAGTCGTCCGAGCCACCACAGAGTCAGCCCTC
 CAGCGGCGCAGAGCGGCGCAGCGCGCACTGGCTGCCCGCGGTAAGCGGAT
 GTGACGTCACTTCGCGGCGCGCTATTGCAACTCCAGCAGCGCCCCGCGGA
 CGCCCCCAATGCCGCGGCCCAAACCGCGCAGCCCCCGCGCGGGGGCGCG
 CCCCCCGCGGCGCCCCCGCCACCCCGCGCGCGGCTCGCGGTGAGTG
 CAGCCCGTAGGAGTGCGGAGTGTTGGGGCGGGGGGGGGGGCGTCTGGAGC
 GGAGCCTTTATCACCGCTGTTTTCCCGATTTCCCGTCTTTTCGCCCCGT
 TTCAGCCCGCGGTACCGGCCCGGTGAGAGGGCGCTGCGGGAGATCCGCC
 GCTATCAGAGCAGCACCGCTCTGCTGCTGCGCGCCAGCCCTTCGCGCGC
 GTGGTAACGGGACTGCCCCGGAACGGGACACCCCCCAACCCCCCAACGG
 GACCATCCCCCACGGATGGATCCCCCCCCACACACATCCAACGTGGGAC
 CCCCCGCCCCAAAATGAGATCTCAACGTGAGATCTGGGGGCCTCAAAATG
 AGACACTCTCCCCCTCCCCCAACGGAACACCCCGAAAATGGGACCACAC
 ATAAAAGTGGGACTCCCCCTCTCCCCCGCCCCGTCAAAATGGAACAC
 CCCCACCTGGACCTTTCAAAAAATAACATTCCCCTCCCCCAAAAATGGG
 ACTTACCACAAAGTGGGATCTTCCCCCAAAATGAACACCCCTCAAAATG
 AGACCCCTCGGACCCCCCCCCAACCCCTCTGCACCCATCNGCCGTCTGTGCA
 CGGAAGGGAAAGGCTGTAGGGTACATCTACCCTTATTTCTTGGGTTTGTG
 TTTTGTGTTTGTGTTATTTAGAAGCAAAACCAAGACAACAAAGCCCAGCC
 AATGCCATTTCTTGGCAGTGGACGCAGGCGCAGGCGGGTTGGTCACAAAG
 CAAGAAGTTGCTGCGGGACTTTGTGCTTTTGGGGCCGTTCTCGTGAACCTT
 CTGAGCCATGGATGAGGAAATTACTTATGCTGATTTAAGGCATCCTACGG
 GCAGTTTGCTCTCTGCTAAGCGGCAGCGCGGTAAGGGATGCTCTGTGTGG
 TGGGTGCTCACCGCAGGCTTGGTTTGGGGGCTTGTGTTCTCTGAGAAAC
 ACCAGCAATGCTGGTTGGGTTCTGGGTCCACCTGGCTTGTATGGGGGAG
 TAAAGGAAGGGGTGGGGGAGAAGGAAGCCTGGGAATGGCCAGAGGTGTGG
 TGGTTTT

FIGURE 10

SUITE 7

2047460-01402

AGAAGAGCCCCGTGATGTCCTCCAGGTGCGGTCCCTCGGTGCCTGTGGGG
ACAACGACAGCCCTAAGCACAGTGTCAACATCCTGGGTGGGGTCCCCAAC
CCAAATCCATGATCTCCCATTTGTCAGGCCATGGTCCTGATGTCCCTCA
GACCCCTCCTAACCATGGTCCCAGCATCCCAATACCTCCACGTGTTTCCAA
TATCCCCACATCCCCCCTCACCAGCCAGGAGCAGTCGGACGGAGACACGC
ATTGGTTTGGCCAGTGCAGTGTGGGTGACAACGCAGCTGTAGATGTCCCC
GTGGTGTGGGGGGCGTGCGGGGATCAGCCGTGCTGCCGCCGTCCGGCTGT
AGGTTCATCGGCTGCCTGGCGGTGACCTGAAGTCCAGCTGTCCATCACT
GTGTCCCTGGGTGACTGTGATGTCCCCGAGCCCCCGGCGCGGCTGCCA
CGTCACCGTCACATCCAAGGGGTAGAAGCCAGACACGTGGCAGCGTAGCT
CTGCTGACGTCCCCGGGGGCCACCACCAGGTTCTTCGGGGACAGCGTCACC
TTGGGGGGCTCTGGGAGACATGTGGGGGGACATCGGTCCCATATAGCCCA
TAGGGCCCCCTCCTATAGGGCTCATCCCCCCTATAAACCTACAGGTGAAC
TATGGGATGATGCCACCCCATCCTATAGTCCTCATAGGAATACCACCCGG
TCCCATCCACCCTATAGCCTCCATAGGAATACCACCCAGTCCCATCCACC
CTACAGCCCCCAGGAATATCACCAGTCCCATCCACCCTACAGCCCC
CATAGGAATACCGCCTGCTCCCATATGTCTATCTGACCAATAGGAATAC
CACCCAGTCATACACACTCCGTAGGAACACTGCCCAACCCACACCCCAT
AGGAACACCGCCTGCCCCACATGGACGCACCAAAGACGTGGAGCTGCAGC
ACTGTCTGTGTGTGCCCCGTGGGGCAGGAACACGGAGCAGATGTAGGTGCC
CTCATCCCCCGGTGATGGCCGCGCCAGCCGAGTGTACCCGCTGTCACCC
CGTCCCCATCCCGTGTCCCAGCAGCAGTTCGGCCCCGGGGGTGGCGCGG
GGGGCGCGGGCGGTGGAAGTGTATAGG

FIGURE 10

SUITE 8

204709150-01402

46/110

AB1B3FOR.txt

CCAACTTCCTTTGGTT CAGGGAAGAAGACTCACCCACTGCTTTGGTTTGT
TGCACTGGAAAAGCATGAAGAAAGCACCACATGATGAGAGGAACAGTTCA
TCCCACAGCTCACGCAGGAAGAACCATTTTTAATTTAATTTGGGAGGGA
GCACTCACCCAGGTCTGAAGCTAGTTTATCTGCAATGAAACAAATAAGAA
ATGCATGATGAGAAGGGTCAGAATATCATCCCATGGCTGATCCCATGGGA
AGACCCCGAATCTCTTTGGTTTGCGGAGGAGGACTCACCCAACGTGTGCAT
TCCTTCCCTCTGCAAAGGGAAAGCAGAAACAGTG

FIGURE 10

SUITE 9

20443160.04402

47/110

AB1C1FOR.txt

TGGGATCAAGTTGAGTAGACATAGCATCCTCGCTTTTAGACAAGACCTGC
ACAGTATACCACCGTTTACTGTGCAGATAATGACCAAAAGCAATATGCGT
CACACTTTTCTGGTGACAACGTCACAAAATGGCGGTCGTCAATCGTGACG
AACAGCACAAACGCCCTTTCTCATCGAAGATTTCATCTGCCAGACCTGG
TGACGCGAACCGAGATGCAACGGTTTGCATACGCCGCGCACCCGCCCTTC
TCGTGCCGAGCGGACGTGGTTAGCATTGATTTCCAGACCAACCACTTTT
GCTCACCTTCGGTACATAAATAACCGGCAACGGAACCGATACTTTCGGCC
ATTACCACGGGTGCTCCTCCATGCAGCAACCCGAAAGGCTGCTTTGTCCG
CGAGTCTACTGGCATTGTGCGTTCAAGGGTGTGATCACC AATATGTTCAA
AGCGAATATCCAGGAACCCCAACCATGTTTCCTTCACCCATAGCATTGAGT
GCTTCCAGGGTGATTTTCCGTTTCCAGATCATTTAATAATCTCCAGTTAA
AGCCTGCACAGGATGGCTTACCCCGTGCCTTCAACCCGTTTTATCTGGCT
ACGGCAAGGAATATCCGGTT

FIGURE 10

SUITE 10

20443150.01402

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AB3A11RE.txt

CCGTCGCCTCGGCTCTCCCTCGGGCTCCACCCCCCGTTCCGCCCCTTTG
CCGCCGCATCTCCCGCTCTGTACCCTCCCCAAGAAGTCGCTCAGACGGCG
TCGCGTTGTCTGCACATCCTCGGGGACCGTCTGTTGTGCGGCAGCAGGGG
AGGGGAGCGGGCGGTCTGTGCTCTTCTATTCCCTTCAGTACAAGAAGGTG
GTTTGGGTTCTTTAACCAAATATACTCTTTTGTTCATATAAATCACC
AGAAGGAATTGGTCTGTTGAATATATAGGAGTGGTGGAGAGAGTCGAAGA
AGTGTTCCTGTGACAAAACACCGTTAAAAGTGAATTCATGGAGAACGCA
CTGCAGTGACACAGAAGGGAAAACACGAAACATAAATAATTTGCCGATTT
ATCATCGATTTCAAGGTCCTTTGGGCTGATTGCTTTCCAGTATTTCCCT
TTGGAGAAAAACCGGTGAAAAATGG

FIGURE 10

SUITE 11

204470-09E400F

49/110

AB586FOR.txt

TCACCTGGCTTTGCTGCTCCAGACCCCGCAGGAAGCGACCCCCCTGGCCC
CTGGCATCCCGCAGCCCCACACGCAGCTGTGCACGGCCCCACACTGGCGC
CCCATCTGGGAATCTGGGGGTCCAAAGGGTCAGTGGAGTCAGGCGGGTCC
AAAGGTCAGTGCGGTCAGGAGGTCCCCAGATGTCAATAGGGTCAGGGGGA
GGGATCCCAAAGGCCAATAAGGTCAAGGGGAGAGATTCCAAAGGTCAGTA
GGGTCAAGGTGCCCCAGAGGTCAATAGGGTTGGGGGAACCCAAAGATTAT
AGGGTCAAGGAGTGACCCCAAAGGACATCAGGGCCACTGATTTGGGGTGG
ATGGGAGAGGAATTTGGGGAGTTCAGGAGAGTTGGAGGGGATTTGGGAGG
TTTTGGAGGAGACAGATGGGGATTTTGGTGGGAATTTGGGGAAGATTGGG
TGGGATTTGGGATTTGGGTGGGATTTAGGTGGGGATTTGGGGGGATTTTG
TCTCTGGGTGTCCCATAC

FIGURE 10

SUITE 12

204110-0140E

50/110

AB6E4FOR.txt

CCTGAAACTTTGGGGTGAGCATCTCCATCAGCTCATCTGCAATGCAATGG
GATCTTCCAGTCTTTGGGTTTTGTGCTCGTTGTGCCACTATTTTCATGGC
ATCCTAAGATGGTGCTGTATTATTTTTGTGACACTGTAAGAGACTGGAGC
AGAAATTTTGTCAAAATTAACAAAAAAAAAAAAAAAAAAAAA

FIGURE 10

SUITE 13

204710-094E400F

51/110

AB6G8REV.txt

GTTCTATGATTTCTTTGGTCCGAATACCATGAAATCTGATAFTTCCATTT
 CAGTATCTGAACTGGGTTCTCTGCTGGATCACAGTGGTCCACACAAAGAA
 GCAGAACAGTATATCGCTCGCGTCTTTAACGCAGACCGCAGCTACATGGT
 GACCAACGGTACTTCCACTGCGAACAAAATTGTTGGTATGTACTCTGCTC
 CAGCAGGCAGCACCATTCTGATTGACCGTAACTGCCACAAATCGCTGACC
 CACCTGATGATGATGAGCGATGTTACGCCAATCTATTTCCGCCCGACCCG
 TAACGCTTACGGTATTCTTGGTGGTATCCCACAGAGTGAATTCCAGCAGC
 CTACCATTGCTAAGCGCGTGAAAGAAACACCAAACGCAACCTGGCCGGTA
 CATGCTGTAATTACCAACTCTACCTATGATGGTCTGCTGTACAACACCGA
 CTTTCATCAAGAAAACACTGGATGTGAAATCCATCCACTTTGACTCCGCGT
 GGGTGCCTTACACCAACTTCTCACCATTACGAAGGTAAATGCGGTATG
 AGCGGTGGCCGTGTAGAAGGGAAGTGATTTACGAAACCCAGTCCACTCA
 CAAACTGCTGGCGGCGTTCTCTCAGGCTTCCATGATCCACGTTTAAGGTG
 ACGTTAAACGAAAGAAACCTTTAACGAAAGCCTACATGATGCACAACAAC
 AACTTCTCCG

FIGURE 10

SUITE 14

204760-04402

CCACCACCGCTTTGGGCGAGTGCCAGTGCTCCTCACAGGCTGTGGGGCAGA
 GCAGGTGACCCCCCAAGGATTTCCCCTACAAAGAGCCCCACAGAGACAGA
 AATCCTTCACCTGAGCTGCAGCAAGCGCGGGCTACACCCAGCATCAATC
 TTTGCCCAGCTTCTACCTTTGCCCAGCTTCTACCTTTGCCCAGCTCCAGG
 GTGCAATGCGAGCAACTTGGCATCAGACCAATACAGTCAAAGGTTGGAGA
 ACATAAAACACATCCCATTGCAGCTTTGTGCACCACCCTGGGTCTCTGCT
 ATCACCAGGAACATGGACACAGGAGAAGCTTTGCCATAGCACAGGAGAAA
 GCTGTGCGCTGCACTTCATGAGCATTTCTCTCAATTTCTCCTGTATCCCA
 CAGGTTACAGGCACCAGTAATTCTGCCAGAGCTATTCTGAAGGGCACGTG
 GTGAAGGATTATGGCTTGGAGCAGTGGGGAGAGCCAAAAGCCCTTCCCAC
 ACTTGATGCACTCCAAGGGTGTGATCCCAGCATGCAGCCTCTCATGTTGG
 AATGGTCAATTTTATCCTAAATCCTCTTGCACTTGGAGCAATGTTGAGT
 TATTTTCCCAGTGTGCATTACAGTGAGGTCCCCCTGAAGCCTACTCTTC
 TCCAGCCAATTTCTTATGATCACGAAGGGGATGATATGATGGTGACATGG
 GGGATTTCCACGTGGATGCTGCAGGGCAGATGGGGAAGGGGTGAGGGGAG
 ATGCCCACCAGCAGAGTTCCCAATCAGGACACAGCAGTTTGTCTGCCAGC
 ACCAGGAAGCAGCTTCCCCTCCTTTCCCTGCTGGGAAATCACTCCTTTGG
 AATGTTTTTTTTTCTGCTGCTCACCCACATTTTGCACAGGGCTGATCT
 TCCAGGTGAGCCCAAATCTGCATCCCCGCACGGATAACCTCTCCCTCCC
 TAAGAATCAGTGCATCCTGCCTGCCTGCAAAAGCAGCTGCTGAGATGTCTT
 TTGCAGCCCTTATTCCCGCAGACCCCGTGCAAGAACACACACATCTCCAT
 CCTCTCCTCCGTTGGCAAGGAATGGGTTTGCAAAGGGATGGGCACAACCA
 GCAATATGCAAAGGAAGAGGTGTGAAAGTCTGGGGAGCAATGAATCTGT
 CCCCCGGAAGATGTTTCCATGGGGCAGTTAAGGAGGAGAATTGGAAATGA
 AGCAGATGATGCAGCAATGAAACTATCCCAGAAAAGGGGGGAAAAGCAAT
 TCTGGTAATGAAGATACATAAAGGAGAAGGGCTTCTCGCTGTCTGGACGC
 AGTTCTGTTGGTTAACGTCTTTTCTCTTTGTGCTCTTTGCACTTTTTTCT
 TTGCCTGCTCTGGTCAGGATGAGGCAGAGCCCTCACGGGGCCCTTTTACA
 CCTTTTTTTAGCACACAGAAGCGCAGCGGCCGTCTCAGCACCCAGCATCG
 ATGAGAAGGACTGCAAATAAATTAATGCGTTACTGAATAGACAGTCGT
 AATTAAGAGTCAAACCCATCCCCTCCCAGTATTCCAGCTGCCGAGGCATC
 GGTTGGCACAGAATCACCAAATATTGCCTTTCTTCCCCCATCCCCGCTTA
 TCAGCCAATGCTCTCTGACCCCTAAAAGGTCTCGATTTGGGGTCTTTTTG
 TTGTTGTTGTTGTTGTTCTGGGTATTTTTAGGCTTTTATTATCAGCGATT
 TTTAGCTTCTCACTGCTTACCCCCCAGCTCAGCACCGCATCGCTCACTG
 CCATCGCTGAACCCAGCGGCGTTTCCATCCCTCAGAGAGCAGCAAAATGA
 GACATCGGCCCGTCTGTCACGGAAGGGAAAGGCTGTAGGGTACATCTACCC
 TTATTTCTTGGGTTTGTGTTTTGTTTGTGTTATTTAGAAAGCAAAACCA
 AGACAACAAAGCCCAGCCAATGCCATTTCTTGGCAGTGGACGCAGGCGCA
 GCGGGTTGGTCACAAAGCAAGAAGTTGCTGCGGGACTTTGTCTGTTTGG
 GGCGTTCTCGTGAACCTCTGAGCCATGGATGAGGAAATTAATTATGCTG
 ATTTAAGGCATCCTACGGGCAGTTTGCTCCTGCTAAGCGGCAGCGCGGT
 AAGGGATGCTCTGTGTGGTGGGTGCTCACCGCAGGCTTGGTTTGGGGGCT
 TGCTGTTCTCTGAGAAACACCAGCAATGCTGGTTGGGTTCTGGGTCCACC
 CTGGCTTGTATGGGGGAGTAAAGGAAGGGGTGGGGGAGAAGGAAGCCTGG
 GAATGGCCAGAGGTGTGGTGGTTTTGAGCAAAAATCAGCCCAGATCGGGA
 AGCCCAATGTGAGAGAATGGAATGAAATGGTGGCAAACGCACCCTGCATC
 CACGTGGCATGAGGGCTGCAGACATCCCCGCCCTCCCAGCCACCGGCTGC
 CCCACACTGGGCTCAGCTCACAAAGCCTGGGGGCTGCTCAGCTTCCACCC
 CATGCTCTATGGAGCCTGCAGGGCCTCCACCACCTCCAGAACCACACGTG
 GAGGTGATGTCCCTGTGTCCATCTGACCTCCAGCGGGAGCCCATCCCATG
 CTCCCTGCTGCTGTCACCCCTCTGTGCCACCTCCTTCCCAGCTGGGAACC
 ACTGGGAGGCCACTGGGAAGGTTCCAGGGGACCCTGGAAGTGGAGGAAAAC
 AAACAGGCATCAACTTCTGCTCATACACAGCATGGGAACCAATGGGAAGG
 GTCCGGGGACCCCAAT

FEUILLE DE REMPLACEMENT (REGLE 26)

FIGURE 10

SUITE 15

10043150-031402

TABLEAU DE REMPLACEMENT (REGLE 26)

SUITE 16

[illegible]

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ACGCGTGGTTCTGGAGGTGGTGGAGGCCCTGCAGGCTCCATAGAGCATGG
GGTGGAAAGCTGAGCAGCCCCCAGGCTTTGTGAGCCGAGCCCAGTGTGGGG
CAGCCGGTGGCTGGGAGGGCGGGGATGTCTGCAGCC

FIGURE 10

SUITE 17

2017-09-28 10:00

CCCAGAACCCCAACCAGCATTGCTGGTGTCTCTCAGAGAACAGCAAGCCCC
CAAACCAAGCCTGCGGTGAGCACCACACAGAGCATCCCTTACCGCG
CTGCCGCTTAGCAGGAGGCAAACTGCCCGTAGGATGCCTTAAATCAGCAT
AAGTAATTTCTCATCCATGGCTCAGAAGTTCACGAGAACGGCCCCAAAA
CGACAAAGTCCCGCAGCAACTTCTTGCTTTGTGACCAACCCGCTGCGCC
TGCGTCCACTGCCAGGAAATGGCATTGGCTGGGCTTTGTTGTCTTGCTTT
TGCTTTTAAATAACAACAAAACAAAACACAAACCCAAAGAAATAAGGGTAG
ATGTACCCTACAGCCTTTCCCTTCCGTGCGCAACGGCCGATGTCTCATTT
TGCTGCTCTCTGAGGGATGGAAACGCCGCTGGGTTCAGCGATGGCAGTGA
GCGACGCGGTGCTGAGCTGGGGGGTAAGCAGTGAGAAGCTGAAAAATCGC
TGATAATAAAAGCCTAAAAATACCCAGAACAACAACAACAACAAAA
GACCCCAAATCGAGACCTTTTAGGGGTGAGAGAGCATTGGCTGATAAGCG
GGGATGGGGGAAGAAAGGCAATATTTGGTGATTCTGTGCCAACCGATGCC
TCGGCAGCTGGAATACTGGGAGGGGATGGGTTTGACTTTTAATTACGGCT
GTCTATTAGTAAGGCATTTAATTTATTTGCAGTCCCTTCTCTCCATGC
TGGGTGCTGAGACGGCCGCTGCGCTTCTGTGTGCTAAAAAAGGTGTGAA
AGGGCCCCGTGAGGGCTCTGCCTCATCCTGACCAGAGCAGGCAAGAAAA
AAGTGCAAAGAGCACAAAGAGAAAAGACGTTAACCAACAGAACTGCGTCC
AGACAGCGAGAAGCCCTTCTCCTTTATGTATCTTCATTACCAGAATTGCT
TTTTCCCCCTTTTCTGGGATAGTTTTATTGCTGCATCATCTGCTTCATT
CCAATTCCCCTCCTTAACTGCCCCATGGAAACATCTTCCGGGGGACAGAT
TCATTGCTCCCCAGACTTTCGACACCTCTTCCCTTGCAATATTGCTGGTTG
TGCCCATCCCTTTGCAAAACCATTCCTTGCCAACGGAGGAGAGGATGGAG
ATGTGTGTGGTTCTGTACGGGGTCTGCAGGAATAAGGGCTGCAAAAGACA
TCTCAGCAGCTGCTTTGCAGGCAGGCAGGATGCACTGATTCTTAGGGAGG
GAGAGGTTATCTGTGCGGGGATGCAGAGTTTGGGCTGACCTGGAAGATCA
GCCCTGTGCAAAATGTGGGTGAGCAGCAGAAAAAATTTTATTTTCAATTC
CAAAGGAGTGATTTCCAGCAGGGAAAGGAGGGGAAGCTGCTTCTGGTG
CTGGCAGCAAAACTGCTGTGTCTCCATGGGAACCTGCTGGTGGGCATC
TCCCCTCACCCCTTCTCATCTGCCCTGCAGCATCCACGTGGAAATCCCC
CCTGAAAAAGCCCATTTTGTGACCATGCATCACATTTATTTTCGCATTCA
GCATCAGACGGACACAGGCAATGGGTGGGGGATGGGGGGGGGTCTGAG
GGTATATCTTTTTGCTGAGCCAGGTTTTGAGTCATGGGGGATAATTTTCA
TCCAAGGGGAGGGGGGCATTTAACTGCAGGTGGTAACAATGAAAGGCAGT
GGGAGTTGTTGTGATTGCATGGGGGAAAGCACTGGTTTTTTTCCATAAATT
GGGACTGATGTGGCTGTTGTTGCTTATTTTATGGGGGAGGGTTGTGGGG
TTTTTTTCCCCTATATTACATTGCATTTAATTTAGTCCTCTCTCATTGT
CTATCCCTGGCAATGCTAGGACTTCTCCTTGCTGTTTTCTGTTGGGCGAT
CATTGCCACAGAGGGAGGAATTGCTTTTCAATTTGGGTCACTGCAATGAGT
TTTAGCACCAGAAATATATCCTTATGGGTCTCTGCTTTTGGGGCACTGC
TGATGGGTGGAAGTTTTGGTTTGCAGGTGAAGTGGAAGCCCCAAATGGA
GGAAGTGAGGGGAATATCCCCATGTTTTGGGCACAGAAATGGAGCAGGAGG
AAGGTAACAGCCGAGCCATGCCCTTAACACATCTGTTTATTGTTATTATT
ATTGTTATTATTTTATTGATTACTTCTTTAACTTGAGAACAAGGGGAGG
GATGTGGGTGGGAAGAAATGAGTCTCATTTCTTTTAGCACTTCCCTCAA
GGGGAAATTTGTGTTGGTTGTTGAGCAGCAGGTGGACTTCTTGCTGTGA
GCAGCCACATTTTGAAGAGTTCTGTTGTTATTAGCATTATTTATGCGAT
TCTGTGATGTTTTTATTATAATTAATTGTAATGAATCCTCCCTGAGGCAC
TGGATGGGGGAAAAAACAACATTTTGGGGTCTACTGCTCACACCTGG
GGTGCATGCTTGCCCATTTGGAGGTCCCTTCTCCCATAGGTCCCAGCCGTG
GGGCATGCGTTACCTTCCAGCTCACGATGGCAGCGGTGTTACAGTGCTG
CTCATCACTGCTGTGCCCTTTGCAGGTGAGTGCTGAGGGTTCCAAAGAGC
AGAGAAAACCCCTTTGGG

FIGURE 10

SUITE 18

10043160 "0.1402

TTCTCCACAGAATCCACCAAACCCACACATTTTCAGGTCCCGTCCAGCT
CCCTGCTCTATGCTTACCTCTTCTGCCTTCTTCCGGAGCACAGCCAGCTG
AGACTGCAGATTTTACACTTCATTTTTGCTTGTGTCCAGTTCCCCTTTT
CTGTGGAAAGCTCATAGCATCGGTCCCCTAAAAGCCTCCAGAACTGGGA
CAGAGCAGGCAGGCAGCAGGGGCTGGAGAGAAAGAGCCGTGAGCATCTTC
AGGTGGGAGAAATCCCACCCAGGAGGATTTCTTGGGAAGGGCATTACCT
GCAGAGCTGTTCCATGTGGATTGGCAGAAGTACTGCTCAATGGAGGTATT
CTCGCAGAGCTCTGTCCCATTCCTCCCGTTGGTCTCAGGGCAGTGCCGGG
CAGCGCTTGGAGGTGGTGTGTTTTCTGAAAGACTTTTGGGCACAACCTG
GGGTGAGACGCGGCCCTATGGGGCCAACCCCGTGGAAACCACGCAGGGTT
GGGGTTGGATCCTCGAGCTCTTTTGCAAAGCCTTTCTGGCTATGGTTGCA
CTCAGTTAATTAAACTGTCTAAAACCATATTTTGTATATAATTAGACATG
ATGTTTACTGCTTCTGTCCCCCCTTGGTTTAAGAGCAGAGAGGCTCTTG
CAGAAGGGGAATTCCTCTCACTGAGTGCCACTTTGGAATTGTTGTGTGATC
ACCCAAACTCCAGTGCAAAGCCCCAGCCCCACTTTGGGCAGAATGAATGT
GTTTTCTGCTCAGAAGAGCTTCGATTTCTGTGCA

FIGURE 10

SUITE 19

2047.0.09E400F

CTGCGCTGGGGATCTTGTCTTTCCCTGGCAATGGGAACAGCTGTTGGGTG
 CCTTTTTTGGGAAAGATCTCTTTATCGGTGCATGAAGAATGAAGCGACTA
 ATGGGGAATGGAAGGAGTGGTGGCTGTTTGAGTAATTGACTGATAGGTTG
 ATGGAGGGATACTTGAATTAAGAGCTTTTGGCTCTTATCTCATTGCCTCT
 GTGCACCAGGTTTGGAGTGGGCCAGGCCCTGGCACGGTCAACTTGCTCAC
 TGTTGGCAATAGGAACATTTTTTGGAGCCTCAGAGAGATTTTGTGGAGGA
 ATGGATGGATCATTGATGCTGCTGGTTTGTCTGGGGGGGACCAATGTGATG
 GATTAATTTTTTTCAGTATAAAAAATAGTTTGTGAGGTGAACTTCTGGTGA
 CTGAGTGGATGGTGGATGGAGGGATGTGAGTTTCTGTGGAGGGATGGAT
 GGTGGAAGGTTTGTGGATGCACTGTTGAGTGTCTGGTGGGATCTACATT
 TGGGGCAATGGATGGATGGACTCTGAGAATATAGACTATAGCTGAGTTGG
 CAATGACCAAGAAGGACCATTGCGTTTTGTTTCTGGCTTCATGTAGGATC
 ACCCAGGAATTAAACCCTATGTCATGGTTTTGTAACCTCGCTATTGGTAT
 TCCACATCATAACATCATGGACAAAAGAGAAGAATAGCAAAGTTACAAAA
 CCATGACACCCTACTTCTGAAAGCAGTTTGAATGCTTGGGGAGCTGAA
 TGTTGATGGTGTGGTGGAGTCGTGGGGGGGAGGTGTCCCTGTGGGGCAG
 TCCCTGGGAAGCTATAGCTATAAGTCACCCCAATGCCCTCTGTGTGGG
 AGTAGTGTGGGTGGGGTCACTGGGATACCACAGTGGGGTGGAGCCCAGG
 GGAGTGTCTTTGAGGTGAGTGGGGGTGAGCAGGGCTCTCTAGAGGCCTT
 TGGGGGGTCCAAAAGGAGTTGATGAGAGAGAGAGTGTGGGAGATCCATGG
 GGGGGCTGCAGGCCTCAGTCCCCTCCATCTCTTGGCAGGTGCCCCAGGAA
 CACTATGGGTGGGGACACTGTGGCCCCGAGTGTCCCTGCATTGGGCA
 CCTCCCCATGTCCCCCTGAAGGCTACAACCTCATCTATGGACCCCCCGG
 TGGCCCCGTGAAGGTAATACCCCATAGCACTCCCTGAACTTCCCAGGGGA
 TCTCCCTGGGTATCTCCTGGGGTACCCCAACCCTCCTTGGGGACCCTGCT
 CCCACCCTGGGGAATCCAAAAGTCCCTCCACCACCCAAGCACCCCTAAGAAC
 CCCACTGCACCCCACTATCCCTTGAGGTCCCCAATACTCCTTTTACAGCA
 TTCCCATCCTCCTCCTTGGCCCCCTTATGCTCTCCAGAGACATTAAACAC
 CCCTGTAATGCCCTTAGGGACCCCTGCAGCAGCCCAATAATCCTCCCAT
 GTCTACCTCCAGACACTGCAGCTGCCCCCTGAAGCAACATCCAAGGAGCT
 GTGGGGCCTGGAGCCCAGTGGACGCTATAGGGTGCAGCTCTGGGGCCGGG
 GGCTGGAGCCCCTTGAGACCACCTTTGACACCCGTGAGCTGGGAAAGGGG
 GTCCTGTGGGGTGGGAAGGGGCACCTTGGGTGGAGGACTCTGGGATACCA
 AATACCTGGATGATTTGGGGTGTCTGGGGACATATGGATGCTGGGTCTGA
 AGTATGGAGGGGGGTACCAAGGAATCTGCATCCTTGGGTGGGGAGCTCTG
 GGGGTTCCCAAGTACCTGAATAATGGGTACCTAGTTAGGGGAATGCCTTG
 GGTGGGGGGGGGGCGGACACAGCGGGATGCCCTCGTCCCTTGGTAGGTG
 AACAGGGACACCCCACTGTTGGGGCCACCTACACTGCTCTGTCTTTCAG
 CACCCCTCCCCACCCACATCCCCGGGACTGCGCTGAGGAGCAGCTCAAT
 GGACCGGGGCCTTCACGAGAGGTCCTCATCTTCTCGGGGGCGACCGGCA
 GCGGCCACTGCACGTCTTCTGCGACATGGAGAGCAATGGGGGCGGCTGGC
 TGGTGGGGAACGGGGCGGTGGGGAGGGTGTCTGGTGGGCTCTAGGGGGT
 GCTATGAGGAGTCTGGTGGGCAATGGGGGTACAGGGTGGGGTGGCTGAC
 TCCATGGTTGCCATTATAAGGGTTGGATTGGCAATAAGAGACCTGTGGAG
 CAACTGGGGGCAATTTGGGGTATCTGGGGAGGTTCTGTGGGGGTTGAGAAG
 CAATGGGGGGGGGAGTGGGGGAGGCTGGAAGATTTAGGGGAGGTTAATG
 GGAAGGTCTTGTGGGGCAATTTGGGGTAATTCTGGGAACTGCAGGGGGATC
 CCAGTGTTCCTGTGAGATTACATAACCCCTATACTATCCATGGGGATCA
 CAGTAACCTCTGGAACATAAATGGGGGAGAAGCCAGGGAGCAATGGGG
 GGCTGTGGTGGATCTGGAGGGGCAATAGGGGTGCCCTGGGGGGCAATATG
 AGGGTCTTAGGGTGCAATGTTGGGGTCTAGGGGGAAGTAATGGGGGGTC
 TGGGGGCAGTGGTGGGGTCTAGAGGGG

FIGURE 10

SUIITE 20

10043160.011402

GGAGGGAGCACTCACCCAGGTCTGAAGCTAGTTTATCTGCAATGAAACAA
ATAAGAAATGCATGATGAGAAGGGTCAGAATATCATCCCATGGCTGATCC
CATGGGAAGACCCCGAATCTCTTTGGTTTGCGGAGGAGGACTCACCCAAC
TGTGCATTCCTTCTCTCTGCAAAGGGAAAGCAGAAACAGTGTGTGGTGAG
AGGAGCAGCTCATCCACACATCGCACAGGAAAACCCCTTTTTTATTTA
ATTTGGAGGGAGGACTCACCCAGTTCTGAAGCTAGTTTCTTTGCTAAAGA
AACAGATAAGAAATGCATGATGAGAAGGATCAAATTATCATCCCATAGGA
ATACCCAGATCTCTTTGGTTAGCGGAGGAAGACTCACCGAACTCTGTGT
TTCTTCTCTCTACAAAAGAAAGGCAGAAACAATGCATGAAGACAGGAGCA
TCTCGTCCCACAGCTCCCAAAGGAAAACCCCTTTTTTGTTTAATTTTAAA
GGCAGCACTCACCCAGATTTTCAACTAGTGTCTCTGCAAAAAGAAATCAAAT
AAGAAATGCGTGATGAGAAGGGTCAGAATATCATCCCATGGCTGATCCCA
TGGGAAGACCTTGAATCTCTTTGGTTTGCGGAGGACTCACCCAACTTTGC
ATCCCTTCTCTCTGCAAAGGAAAAGCAGAAAGCAGTGCGTGATGAACTGAA
CAGCTCATCCACAGCTCACACAGGCATCCCTCATTTTGTATTTTGTGTG
GGAGGGAGGACTTACCCAGTTCTGCAGCTAGTGTCCCTGATAAAGAATCA
AATAAGAAACGCATGACGAGAAGGCTCAGGTTATCATCCCATGGCTGATC
CCATGGGAAGTCCCCAAATCTCTTTGGTTTGAGGAGGGAGACTCACCCAA
CTTTGCATCCATTCCCTCTGCAAAGGAAAAGCAGAAACAATGCATTATGA
GATGAATGACTAATTGCACAGCTCCCAAACATTAAAAAAAAAAAAAATAG
TGGGAAGGGAAACTCATCCACTATCGCAGGTAGTTCTGCTGGAAAAGAAA
GAGCAGAGCAGTGCATGGTCAGAGAGGACAGCTGCTCATCCACAGCTGA
TGCCATGGGGAGACCCTGAATTCCTCACTTTGGGGAAGGAGACTTACCC
AACTCTGCATCTTTTCCCTCTGCAAAATAGAAGCAAAGGAAATGCATGGT
CAGAGGGAACACCTTCTCATCCCATGGTTGCTCCCATGCCAATACCCCA
AATCTTTGTTCTGGTAAG

FIGURE 10

SUITE 21

20443160-011402

CAGTGACAGTGCAGGTGGAGGGACACGGCGAAGGGACGCTGACGGTGGGT
GGCTGCATGGACATTGGTGTCTCCAAGACCGATGTCCCCTCACAACC
TCCCCTCATGGTGTCCCCTCATGCTGCCACGGTGTCCCCTGCTGTCCCAT
CATGGTGTACGCTGTCCCCAGGTGCTCCGCCAGTTCCGCCTGCTGTAC
CTCCGAACGCCACGTGCCAGGCGCTGCACCTGGAGGTGGCCATCACCGGC
CCCATCCTGTACCATGGTGAAGCCCCACCCAAAGGCCCCGCCCCCTTTTC
CTCGCGGGGGCGTGCCTCAACCCTGTTTTGCATATCCCAACCCCCAGC
AGATGAGGACTACGAGGACTACGAGGACTACGAGGAGGCGGAGCCTAAGG
AGGGGGAGGAGCCTACGGAAGGGGCAGTGCCCGTGGAAGGGGCGGGGCCA
GCAGATGACCCCGCCCCCTCAGCCCCGTGTCCTTATGGGATGCCCCGTAA
GCGGCAACGCCGCAGCACACATAACCCTGCCACGAGGTGGCCTTCCTGG
TCTGCTTCCGGTGAGGGGCGGAACCTCCTGTCCCTGGGGGCGGGTCTTCC
TGCTGATGGGCGTGGCCTGTTGTAGGCGGAGCCCAGGGGTGGCACTGACT
GGGATGGCGGTGGTGGAGTCACTCTGCTCAGTGGCTTCTCACCCCATAG
AGCTGACCTGGACAAGGTAGGGGCCAGGGGGACTTGTGGGACATGTTGG
GGGGTTGAGGGGAGTTATGGGGTGTGGGGTTTGGGGGTGTTGGAGTTGTT
GAGGTGGCAGAATGTTTGGGTGGAGTCATGGGATATGGGG

FIGURE 10

SUITE 22

10043160 011402

CCACTCTTGGGTGAGCTGACAGCGTCCCACGTGAGCCCCGACTCCGTCCA
 GCTGGAATGGAGCGTCCCCGAGGGCTCCTTTGACTCCTTCACGGTGCAGT
 ACAAGGATGCACAAGGCCAGCCACAGGTGGTGCCCGTGGACGGTGGGTTG
 CGCACAGTGACCGTGCCCGGGCTGTGCGCGTCCCGCCGCTACAAGTTCAA
 CCTGTATGGGGTGTGGGGGCGGAAGCGTCTGGGCCCCATGTCCACTGATG
 CTGTACAGGTGAGCATGCTGTGTTCTGCCTCCATGTTCTTTTGCTTTCA
 GTGTAGTTGTCATGTGGCAGGAACCTTTGAGGGCCACTTTTGGTTAATGT
 TGCCTTAATAGTCAAGGAAACAATTTGTTCTTGTTGAGTGGGAATGCCTA
 ACGGGATGGGAGTTTGGATGATGAGAGGACAAATCTTATAAGGGATGATT
 GATAATTATTGCGGAACGGATGGAAGGAAGGTTGGATGGATGGAATGGTG
 TTTGGATAAATTTGTGCTCAGAGCACAGCTGGAGTGGTGGATGAATGTTG
 CTTTGCTTGTTGAATAGATGGATGTTTGGTTGTGTGGTTGCTTCCACTGA
 GAATTCCTCCCTCTGTGCTGCAGCAGCAGCTCCAGCACAGAGGAGCCAC
 CTTCCCCACCACGTCTGGGTGAGCTGACAGCGTCCCATGTGCGCCCCGAC
 TCCGTCCAGTCAAGGATGGAGCGTCCCCGAGGGCTCCTTTGACTCCTTCAC
 GGTGCAGTACAAGGATGCACAAGGCCAGCCACAGGTGGTGCCCGTGGACG
 GTGGGTTGCGCACAGTGACCGTGCCCGGGCTGTGCGCGTCCCGCCGCTAC
 AAGTTCAACCTGTATGGGGTGTGGGGGCGGAAGCGTCTGGGCCCCATGTC
 CACTGATGCTGTACAGGTGAGGGCAGGAATTGGCACCTGGTGGGCTCTG
 GGTGTCAGCAGGTAGAAATGTAAACGTGGCCTGCGCTGGGGATCTTGTT
 TTCCCTGGCAATGGGAACAGCTGTTGGGTGCCTTTTTTGGGAAGGATCC
 CTTAATCGCAGCATGAAGTATGAATGGACCAATTGGGTGTGGGTGGAGTG
 ATGGCTGTTGAGATGAGTTGGTGGCTGCTTGAGTAATTGTCTGTTGGAAT
 GGATGGACAGATATGTGAAGGAGTGAAGGATGGATAAAGTAATTTAGGA
 ATCGGTGGATGAAGAATGGGTAGGTAGACCCTTGGTGAAGTGGTAGAATG
 GAAGGATTTATGAACAGATATGAGTTAATTCTTGCAATCGAAGTAGGTGTA
 AGTGTCTATTAGCCTGTTGCACTGAACATGCAGTTGCATAGACAAATGAG
 TGGGGAGAAGTACGGAGTAAATCCCTGCATGAATGGTAGGACAGAAACCT
 GAATGCCTGGATGCTGGCAGTGTGAAGAATGGCACTTGGGATAGATGGTT
 CGAGTATGGGGTAGATTAAAAGATGGATGGAAAAGAGGAACAGAGAGAGG
 GTGATTGGATGAATGGATGGATGGTTGGATGTGACTGATTGACAGGTACC
 AAGCTTTTTTCTGCACTGTGCCTTCTGTGCTGCAGCTGCAGAAGAGACG
 GAGGAGGAACACCGTCCCAGCCACGCCTAGGAGAGCTGACGGCATCCCA
 TGTGAGCCCCAACTCCGTCCAGCTGGAATGGAGCATCCCTGAGGGCTCCT
 TTGACTCCTTCACGGTGCAGTACATAGACGTGCAAGGCCAGCCGAGGAG
 CTGCACCTGGATAGTGGGTGCGGCACAGTGACCGTGTCTGGTTTGCTGCC
 ATCC

FIGURE 10

SUITE 23

20443160-01402

GCACAGAAGGAACCGCCATCCCAACCACGCCTGGGTGAGCTGACGGCCTC
 CCACGTGAGCCCCGACTCCGTCCAGCTGGAATGGAGCGTCCCCGAGGGCT
 CCTTTGACTCCTTCACGGTGCAGTACAAGGATGCACAAGGCCAGCCACAG
 GTGGTGCCCGTGGACGGTGGGTGCGCACAGTGACCGTGCCCGGGCTGTC
 GCCGTCCCGCCGCTACAAGTTCAACCTGTATGGGGTGTGGGGGCGGAAGC
 GTCTGGGCCCCATGTCCACTGATGCTGTACAGGTGAGCATGCTGTGTTT
 TGCCTCCATGTTCTTTTGCTTTTCACTGTAGTTGTCATGTGGCAGGAACCT
 TTCAGGGCCACTTTTGGTTAATGTTGCCTTAATAGTCAAGGAAACAATTT
 GTTCTTGTTGAGTGGGAATGCCTAACGGGATGGGAGTTTGGATGATGAGA
 GGACAAATCTTATAAGGGATGATTGATAATTATTGCGGAACGGATGGAAG
 GAAGGTTGGATGGATGGAATGGTGTGTTGGATAAATTTGTGCTCAGAGCAC
 AGCTGGAGTGTTGGATGAATGTTGCTTTGCTTGTGTAATAGATGGATGTT
 TGGTTGTATGGTTGCTTCCACTGAGAATTCCTCCCTCTGTGCTGCAGCAG
 CAGCTCCAGCACAAAGAGGAGCCACCTTCCCCACCACGTCTGGGTGAGCTG
 ACAGCGTCCCATGTGCGCCCCGACTCCGTCCAGCTGGAATGGAGCGTCCC
 CGAGGGCTCCTTTGACTCCTTCACGGTGCAGTACAAGGATGCACAAGGCC
 AGCCACAGGTGGTGCCCGTGGACGGTGGGTGCGCACAGTGACCGTGCCC
 GGGCTGTGCGCGTCCCGCCGCTACAAGTTCAACCTGTATGGGGTGTGGGG
 GCGGAAGCGTCTGGGCCCCATGTCCACTGATGCTGTACAGGTGAGGGCA
 GGAATTGGCACCTGTTGGGCTCTGGGTTTGCAGCAGGTAGAAATGTAAAC
 GTGGCCTGCGCTGGGGATCTTGTGTTTCCCCTGGCAATGGGAACAGCTGTT
 GGGTGCCTTTTTTGGGAAGGATCCCTTAATCGCAGCATGAAGTATGAATG
 GACCAATTGGGTGTGGGTGGAGTGATGGCTGTTGAGATGAGTTGGT

FIGURE 10

SUITE 24

2004-10-01 14:00

CTGTGTCCCCAACCTGCTTGCTTGGTGTTCCTCGCAGGACACGCTGGTGGCCCTG
GAGGCGCTGGCCCAGATGTGGCTGCACTGGGGCCGTGGGAACACAATGGG
GCTGAACCTGGGGCTCTCCTGGCCGGGGGTGCCGGGGGAGGGCTGGTG
GCACTCAGGTTATGCTGAAGCCGGGGCTGGAGCCGCTGGAGCAGGAGCTG
CAGGTGGGGACATGGCGGGATGTGGGGACACGAGGGATGTGAGGACACTG
GGGACATGTCTGGACTTGGTAGGATGTAACATGAAGACACTGGGGACATG
GTAGGACATGGGGGACATGAGAACACGGGATGTGGGGGACATGGTAGGAC
ATGATGGACACAGGGGCTTTGGGGTCCTTGGGTCTCGCTCTGTCCCCATG
TCCCCAGGTGCCTCTGGGCAGCCAGTGACAGTGACAGGTGGAGGGACACG
GCGAAGGGACGCTGACGGTGGGTGGCTGCATGGACATTGGTGTCTCTCC
AAGACCGATGTCCCCTCACAACTCCCCTCATGGTGTCCCCTCATGCTGC
CACGGTGTCCCCTGCTGTCCCATCATGGTGTCAAGCTGTCCCCAGGTGCT
CCGCCAGTTCGGCCTGCTGTACCTCCGAAGCCACGTGCCAGGCGCTGC
ACCTGGAGGTGGCCATCACCGGCCCCATCCTGTACCATGGTGAGGCCCCG
CCCCCTTTTCTCGCGGGGGGGCGTGCCCTCAACCTGTTTTGCATATCCC
AACCCCCAGCAGATGAGGACTACGAGGACTACGAGGACTACGAGGAGGCG
GAGCCTAAGGAGGGGGAGGAGCCTACGGAAGGGGCAGTGCCCGTGGAAGG
GGCGGGGCCAGCAGATGACCCCGCCCCCTCAGCCCCGTGTCCTTATGGG
ATGCCCCGTAAGCGGCAACGCCGCAGCACACATAACCTGCCCCACGAGGTG
GCCTTCCTGGTCTGCTTCCGGTGAGGGGGCGGAACCTTCTGTCCCTGGGGG
CGGGTCTTCCTGCTGATGGGCGTGGCTTATTGCTGAGGGGCG

FIGURE 10

SUITE 25

CCTCTGCTGCTTCCAGAGCAAAGGAAAAGGGAGAGGGGGGCTCCCACCAC
CCTATCCCAGAGCATCAGATGGGCAATGGATGCAGCAGCTCCGTGGGTG
TGGAGGTGGCACGTGGCAGGAGCGAGGACGGCTCGGAGATACCGAGGTCA
TCAGCCACCGAAACCATCTCAGGAAAGGAATTTCCACACAAAACCTCCAT
TTGGAGCACCTGGCAGAGAAGCTGAAGCTTTTGGAGCTGGATGGAGACAG
AGGGGAGAAGGAGAAACTCTGCTCGTGGCGCAAGAGGACATTCCCCTCCA
ATGGACCACGGGATGATGGAGGTCCCCTGGAGCCCCCATAAAGGAGTCA
GTGCAGGAGGATGTGGTCAGCCCTGTGTTATTCCCTAAAGCCCTGTTTAA
TCCTTCATGTCCATGCTGAAAACCTTCTTCTGCGAAGTCCAACACATTG
CATCTCTTCCCTTCTTTCTCCCATCACAATATCCTCCCCAAACCCCTTTT
TCTTCCTCCAGGAGCAGATTACAGCGATCTGGAGAACCTCAAGAAACAA
AAGGAGGAGCTCTTAGAACTCAAAGGAGTGGGGAGAGGCGATGCCAAGA
CCTTCTGGTAAGAAGCTGTTGCCTTCAAGCTGGAAAAACAGAGGTCTTTT
TGGGGTCCACGTTGTTGATTTTCCACAACCTACAGACACGGACGGAGGCT
GAGAGGCAGAAAATTGTGTCAGAATTCCGTGAGCTCCGCCGTTTTCTGAA
GGAGAAGGAGATGGTGCTCGTGGCACGGCTGGGGGAGCTGGACAGGGCTG
TGCTGAGGAGGCAGGAGGAGGAGGAG

FIGURE 10
SUIITE 26

10043150 01402

AGCCCAGCACTCTGCAGTCTTCTATCAGTTCCAATAGAGGAATTTTGGTG
GTAGAAGGGGCTGGAAGGACTCACTCTGCTTTGTGGTCTCAGCTGCTGGA
AAACAAAGCAGAGAAATAGCTGGTCAGCAGGGCAGCTTGGTTTCTGGGGA
CGTCTCCAGAGGCTCTGGACCTTTCCACCTGCCCCACGGTCCACCCACAT
TCCTATCTTTCCGCCCCACACCCCTTTTTCCCTTTCCTTCATTCCCAATCA
AACGGCAAATGTTATTTAATGACCACTGTCAATCCCCAGAAAAATCTCCC
TTTCTCCTGCATACCTCCACGGACCTGAGCTCAGCACCACCCCGACCATC
CCTATCCCTGCTCAACACCTCCCTGTGATCCATCCCCTCCATGCTCAACT
CACCTTTCTTCCTATAGAGAAAAACAGTGATGACAAATGACCCAACCAGA
ATTGTGACGATCACAGCCAGAGCCACCTTCCAGGGATGGGTGATCTGGGA
AAAGGGGTCTGGAAAAACATCAGGACAAGGGTTCTTTTCCATTCCCAT
AAGTGGAAAAGCAAGACTCAGCCTTGGGACATCACAGAACCCAAAGGGGC
AGCAACCAGGGAGCAGTGATGCACAATGACGGCATCCCCATATTGGCACA
GGTGGAGGAGCTGCTCAGCATCGTGTGCCCCACTGCCACTGAGCCATGGAG
AAACCCATCCCAGAAATCCAACCCAACCACCTCATCCATGCAGACTTATC
CACAAATTGCACTGTGCACCTGCTCCAACACCAGCATCTCATGGAACAAT
TTAGCTCCGACCTCTTCCAAAGGCTGCTGTCTTTCAGCTTTCCATCCATG
GATGTGAGGATGAGGATGGACAGAGGTCGGGGTGGGACACACAAACCCAG
CAACACCTGGAGGCGTCACCCCAGCCACTGACCTGACACCTCCAGGTCCA
CCACAGCGTCTGCA

FIGURE 10

SUITE 27

204770-09T400T

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Contig99.txt

CCCAGCAAGGCCAAGCGCCGCCATAACGTCAGTGCCGGTGAGACTGTCTG
ATGCGGTTGCGCGAGGAGAGTCACTGAACATCGGTGATTTAGGCGCAAAG
TATTTAGCGATTGATTGAGGTTCAATTATGCGGCTTCCTTCTGTGGCTGG
TGGGTTTTGGTCTGGCTGTGCTTTACTATTGGCGGCATGCTGGCGCGCTT
AACGCTTTCGGCCTGGTATCGGGTTATCTCGTCTCTGGTCATGATGGCCT
CCGATTCCAGGCGCGAATTGCATCGCGCTTTGTTGGATAGGTGTCAGTTA
TCGGCTTAATCAAGCATTGCTTTGTTGAACAACCGGCGTAGACACCATCA
CCATCAGAAAAAAGTTCTGCGCCGCCGCCACAGAACGGACACTCAAGCAG
AAAAGCCCCAATGAGGTAGCTTGAGATCGAATATCATTGGTTTTCATGCTGC
CTCCCGCTGTTTTAGTGCTTTGAGCTTGTCGCGGTAATCATCCCGGATCC
GGATGAAGTCTTCACGGCGGTAGTTGGTCATTTTCGTGGGGACCATTGAGC
CAGTTGACGTATCCCTGACCGTAACGAGCGACCAACCCAGCTTCGTATTG
CTGCGCCACGGTCGCCTCTTTGGCGGTGTACTTGCCAGCTCCGGCATTAC
AGGATTTGCACTGCTTATGGGCGTTGCGTTCTTCAAAGCGCAGTTCAGGG
TAAGCACCTACTGTCTTGAAATGGCCGCAATCCCACTGGCCACCATGCAG
ATCAGGCGGATTGGTCTCGCCGCGAGCTGATGCATGGCAAATCGGCGTCGC
GCGCACGGATAAAGGCGTTGAAAGCTTTCTGAGCCTGAGCCTTGTAATAT
CCGTCTGGCCTGAGCTCTGCCAGCCGCTCCTTGCGGCGTTTGCGCCCGTC
CTTTTCAGCCTCTTTTTGCTCCTTGATGCGCTTAGCCGCGGCTTTACCT
TCTCCTTCTTGCGTTCTTCCATTGCGAGGATTGCGCCATGCTCCGGGG

FIGURE 10

SUITE 28

10043160-034402

CCCTATGGGGCCAACCCCGTGGAAACCACGCAGGGTTGGGGTTGGATCCT
CGAGCTCTTTTGCAAAGCCTTTCTGGCTATGGTTGCACTCAGTTAATTAA
ACTGTCTAAAACCATATTTTGTATATAATTAGACATGATGTTTACTGCTT
CTGTCCCCCCTTGGTTTAAAGAGCAGAGAGGCTCTTGCAGAAGGGAATTC
CTCTCACTGAGTGCCACTTTGGAATTGTTGTGTGATCACCCTAACTCCAG
TGCAAAGCCCCAGCCCCACTTTGGGCAGAATGAATGTGTTTTCTGCTCAG
AAGAGCTTCGATTTTCTGTGCAGCAATGTGGTTGGGATCTGATCACTCAC
CGCACACGCTGAGCCCTGTCAACCAGCAGCAGCAGCAGCAGCAGCAGCACC
CCCAGCATGCAGGCTTTCTGGAAGTCCCACGGAAGTGGAAAGAGCCCACAC
TTATATAAAACAGACATTTTGAAAAAACTTTTCTTTTACAGAAATGATC
TCCCTGTGAAAGAGCCCCTCCACCAACCTGCTACGTTAGAGCAGAAGTTG
ATGGCTGCTTTGGTTCCTTGAGAATTTGGGGTCCCCGGACCCTTCCCATT
GGTTCCCATGCTGTGTATGAGCAGAAGTTGATGCCTGTTTGTTCCTCC
AGTTCCGGGGTCCCCTGGACCCTTCCCAGTGGCTCCCAGTGGTTCCCAGC
TGGGAAGGAGGTGGCACAGAGGGGTGACAGCAGCAGGGAGCATGGGATGG
GCTCCCGCTGGAGGTCAGATGGACACAGGGACATCACCTCCACGCGTGGT
TCTGGAGGTGGTGGAGGCCCTGCAGGCTCCATAGAGCATGGGGTGGAAAGC
TGAGCAGCCCCCAGGCTTTGTGAGCCGAGCCCAAGTGTGGGGCAGCCGGTG
GCTGGGAGGGCGGGGATGTCTGCAGCCCTCATGCCACGTGGATGCAGGGT
GCGTTTGCCACCATTTCATTCCATTCTCTCACATTGGGCTTCCCGATCTG
GGCTGATTTTGTCTAAAACCACCACACCTCTGGCCATTCCCAGGCTTCC
TTCTCCCCCACCCTTCTTTACTCCCCCATACAAGCCAGGGTGGACCCA
GAACCCAACCAGCATTGCTGGTGTTCCTCA

FIGURE 10

SUITE 29

CCGGCATCACCGGCGCCACAGGTGCGGTTGCTGGCGCCTATATCGCCGAC
 ATCACCGATGGGGAAGATCGGGCTCGCCACTTCGGGCTCATGAGCGCTTG
 TTTCCGGCTGGGTATGGTGGCAGGCCCGTGGCCGGGGGACTGTTGGGCG
 CCATCTCCTTGTCATGCACCATTCTTTCGGCGGGCGGTGCTCAACGGCCTC
 AACCTACTACTGGGCTGCTTCCTAATGCAGGAGTCGCATAAGGGCATCGG
 TCGACGGGATCACGTTGTGTCCCTGAAGCTCTCCTGTACCCAAACACAAA
 GGTGATGTCCCCAGCATCCCTATCCCAGCACTCTGGGGGACTCCTATTGA
 ATTCTCCTTGGGCTTGCTGCCTTCTCTTCCCGTTCCCAGAGATCCCAAA
 AGGTTAAGCACCTTTGGGTGAGTGTTCAGAATTGTCACTGCCAGTTTGG
 GGTATCAGTGGCAAATTGAGACCCCTTTTACCCAATCTTGCACCACTCTGG
 TTCCCCAGTCTTATGGTTTTAGATGGAGTAAAAGGTTTTATATGTCATAA
 AGTTCTTCTGTGTCTGGTTATTGCTGCTTCTGGATGCCAGGATCATGGG
 GATAAGGGGAAAACAATGGGTTCTCTTATGCGTAGAGATGCAATCAGATG
 GGGAGAAAAAGAAATCTTAATCTTCTGATCCATCTGACAGATATTGAGT
 ACAGCCCTGAGGATGTGGGAAATAAATCTGAAGAGTTTGTGGCAGTTCC
 AAGGATTTGGAATGACTAAATCCCATTCCTGGTGTCTGCACAAAAGTTGGC
 TGTGTTGGAACCCAGAAAGATCCATGCAAGTGGGTGATCCCTGAAAGCAT
 TGTGTTCTGCTGTCTGCTAGCGGAGAGAAAAGACACAGAGGGGAAAAATTAA
 GTGTTTTATTGTTAATTATTGTACACTCTGAGGTTTCAAATACCAAATCT
 TTAACGAGAGCGGACCCTTGATTTGAGGGTGACCATCTCAGATGGGGAC
 AACTGTACCTGATCAGGCAAACCTGGGGGAAATTTGCCTTTCTGCCACTC
 TTTTGGGTGGGATTTTCCCTTTTGACCAACATTTCTACATTCTAATCAC
 CCATTGCAGCACTTCTCCCCCTTTTTTTTGGCCCATTTTTCTCCTGCTCA
 GCACTTCTTAACAATATAATATAAATCAATATCATATCAATATGATTCTA
 TGCCAATAGATTAATGGGGATGAAAGACACATAAAAACCCAAGTCTCAT
 TTCATCTGCTTCCCATGGGATGGGTGGGGAGGTGGCTGTCCCCTGAGGCT
 GTAGGATGTGGGGTCAACCTTGTCTGTGTCTCAGGGACACAGCCTCAGCT
 TGGACCTGACCCCTACCACCCACAGCCACGGACGGACCCCTCTCCCCAGAG
 AAGGATGCATGGGAAAAAACAAGATGAGCCCCCTTCATCAGCATCAAA
 AAATGCCACCGTCCCTCCAGCGTAGTCCAAGTGGACGCTGACCCTCCTGG
 GCACCCAGCGCAGAGCTAACAGGGTCACTTGTGGGTGGTGAGTGCCCGG
 ACCTGTCCCCCCCCATTTCTCCACCCCCCAAATCCCCCTTTGGGACAGAG
 GCTGAGTTGACCCCTTCCGAGGGATGGATTCTCGGGCCACACCGATGGCCC
 AGTCCCCTTCATCCCCCACTTCCACCTCCCAGCAGTGCCGGCCGGCAGAG
 AAGCTTTGGTGGCCCAAAACAAAGGGCCAGTAGGCGAATCTTTCGGGGTT
 ATCAGGAAGGTCTGTGTCTTCCCCACGTTTCACTCTTTTCGGTCTT
 CGGAGAGGATGAGGTGAGGCTGAGCGGTGTGGGGTCCAGGCTGATGCTG
 GCTGTGGGGTGGAGAGGATGAGGAGTGAAGGTTTGGGTCTCGGTGCTG
 AGGCCATGAGGATGCGGAGAGCTTGGATCTCCAGCACTAAAGGAGTTGGA
 TGTGCTCTAGATGGCCCCACCTGAGTAGGGTTGTAGGGTGGGACCGTCCC
 TTCCAACCTCAGCCATTCTGTGGGGCCATGGGTGGCATCGGAAGGGTAA
 AAAGTACCAAAGAAGAAAGTAAAAAGGTGAGAGGTGGAACCCCTCTCAT
 GTGCCCGTGCTATATGACAATAAAAGTGTGTTGAGCCCCCAGAATGCCCA
 GAAATAAAGGCGTTTCTGCAGACCTTCTGTTCCATTGGTCAAAAGAAATG
 GTGAGGGGAATAAAAAATGGAAGGAAGGAGATCTATGGGATATTACCTGCA
 AAGTCTGCAGTGCTTCATCTCCTAGACCAACCCGGACCAGTTTCAGCCAAC
 CCCATGGTTTTAAAAAACAGAGCTGAAATCTGAAGGCAGGGATAATGAATG
 AGTCAACCCGCTCACCATATTTGTTTATGGGAAATGGATATTTATCAAG
 GCGAGGGATCTGCCCTGGGGCCATCATCCCAAATTACAGCCAGACTCGGC
 CTGCAGGGTGAAGAAAACCTTGTGTTGGCTGCCCTGATTTTTGTGTATTCCT
 CCTCGGCATCTATTTTTGTCCATTTGGGTACAGCCTATGGGTCCAGGCG
 CGCCTCCATCTAACAGGTAATGCGGCTTTAGGTTCTCATGCTCAGCAAAA
 GGCACTTTTAGGAAAGGTGAAGCTGGAGGGGTGCAGAGCCGGAGAGCAGC
 CCGTCTTACCCCTGAGCACTTCTCAGGAATTACAGCAAAACGTGTAAT
 TAAGASTGGCAAACGGGTATCGAGTCTTTCGGTCTCAATTATTTTCT

FIGURE 10

SUIITE 30

10043460 "011402

GAGTGGGAATAACCCGTTGCTCTTCCATCTCTCTGCATTATTCTGCTGCA
 GAACGAGTGATGGGCTGCTGGTTTTTACCAAAAATACCACCATTTCACC
 CGAAACCCCTTCTGAGTACCTTGAAGCCTCTTCAGGGTTTTCTTCAGAGCA
 CCGTTCCCTCCATGAGGAATGGCACAGCCTCTCCTCCGGGCCCTGGAGAAGC
 GCCCGCTGGCAGCTGGAAGGTCACTTTTTCCACACCTGGAGGGGAAATAAA
 TGCATTTTTAGGTGGTTGTATCACAGAGCATGCCATCACTTCAGGACAGC
 AGAGGCCAGCACACGGCGGCCATCCCCAAAATACCCTTCAGGGCTCGCAG
 TTCCCCTGGAGCAGAAGAGCATTATTGATGAGCTTTCTCCTCCATGGTC
 ACTGCCTGATGCAAAGCTCACAGAACAGCTTTTCAGAGAGGCCACATACC
 TGGTGATGGGGCTTTTACATCCTGGGGACAGAAGAGAGGAGGGGGAGAG
 GAAACTCAGGTGAGTGCATGACCCATTTTGTCTTTAAAGTATGGAAAATT
 GAGCTGTTTGAAGTGGGGGTGGACCTCTTGGGTCTTCCAACATGTGCCCAA
 TTTTGACTTTAAGTCATAGAAAAAGTGAATTGTTTGAAGTGGGGATGGATC
 TGTGGGTCTTTCAACACATGGTCCATTTTGTCTTTAAATCATAGAAAAT
 AAGAATTGTTTGAACAGAGATGGACCTCTGGGGTCTTCCCTCCACGAGGAA
 GGTGAACCAACTGAGGAGCATCCATGCACGGCAATGAATCCTGCAGATCC
 ACCCCACTGCTGCTCTCCCAACCCAGCCGTGGATTTCCTCTTAAACA
 GACCCCATGAGGACCTTCTGCAGTAAGGTGAAAATACTGGGAATACTGAG
 ATGAGGATAAAACGGTGGGGGGAAAGAGGAGGCTGCAAACCTCCATCTCC
 TCATTGTGGTGGGGGTTTTAGGCTGATGGAACGGGCATAAAATGGGAGGAA
 AACACCCAATTAAGGCACCATGCAATTGGTGGGGGTGGGGAGGACATCCC
 TAAAGGACTTTTCCCCTTGAAAAAGCTTCCCTGGAGGAATTCACTCACCG
 ACTGCTGGCTCTTCTCTCCCTGTGCTTTCGTATCCAGCGGGGAAATCTCC
 TCCGAGTGCTTGGCGGTGCTTTTCTGCCTCTTCTCAATCTCATTTTTAG
 GTCTTCCAGCTGCCAGAGCAAGAAGGGCTCTGTGTTTTCTGCCTGGAAT
 CTGAGCCCTCCCTACTGGGGCTCAGCTTTCTCTGATGCAGAAAGTGA
 AAATAAAGAGCAGTGGGACTGGAATAACAGGGGGGACTCATGAGTGGCA
 TCCCCCACTGGAGGAGCTCAATGGTGAGCTGGAATCCTTGCTAAGTTTTA
 TCGAATGTGGGGGACAGGAGGAAGAAATCAAACCTCAAAAAGTCATGAACA
 GGTGGCTGTGAATTGGGGGACAGAAAGCTGAGGGCCCTAAAAGCACAGGAG
 GCAAAAAGGATGGAGAGAAACGACCCTACTGATGACACATCGCTGCCAG
 CAGCTGACACCTACCAGATCCTCCAGGTTTGGGCACTCCAGGGCGCTCTT
 CTTCTCGGAGACTTTCTCTCTCCTCCTTTGGAACCCCTGATATCCCTC
 TGAGTTTCTTCCCAGTGAACCCACAGAACCTGTTGTTTTAGCCCTTTG
 ATGGGGTTGGGGTTTTCCCTTCTGTTCTTCCAGTCTGGGGTAGAGCT
 ATGGGATGGCTGCGTTGAGCCTGCAGGTCTGCTCCTGGTGGCACCCCTGG
 CAGGGCGTGCTGGGAGCTCTGGGTTTGTCTTTGTCTTTCTCCAGTTCC
 TTGTCCCGGGGAGATGCTGAACAATGTCACTTTGCAGATTTTGTGAGCTT
 CCTTTTAGGATCGAGCCATCGGGAGTGGGGTTAGGGGGTGTATATGGGGA
 AACCATAAGGAAATAGGGAAGGAGATGCACAGCCGATCCTTGTGGGGAT
 GTGGAGGAGCACAAGTGAGGATCTTTGGGATTTGAGTGCTCTCTCAGCCC
 AGCACTAACACAGAGCACTCACAGCCCTGGCTCTGAGCTCTCGAGGAAAC
 ATTTCCAACCATTTCTGCCCCACTGTCCTTGTGTTGAGCCCCATGGCCAA
 ATACACATGCCTAGAAAATAAAGCCATGCATTACATATGTATTTAATTTT
 TGCGTGGCAACCACTGAGACCCAACTGGAGGAGATAACTGCCATTCACTT
 GGGCAGTTTGCAGGGGTGAAGTGCAGTTCCAGCAAACCCCTCCCTGTTGG
 GAAGAGCCACAGGGATGGATGGCACTCTGGGAGCTGAAGAAGTGAAGCA
 AACTCCCTGCAACCGCTCCCCTGGGGCACAGAGCCTTTTCATCCCAAAATA
 AGGCGTCCATCATTGAGCAAATGAGTCAACCGTTGGGCAAACGACTTGC
 ATTGCATCCCAGAAAGCATTAAATTGCAGAGCCTGGAAAAGTCTGGGCT
 GGAAACATCTGCATTGCAGATCTATGGAGCAGAAAGACCCCTGAACAGAT
 CCTTCACCCAAATTTCCCAGCAGGTGGGACCAAATGGCAGCGATGCGTGG
 GGCTGAGGAAAGATACCAACACATCAAAGAGCAATATTGAAATTTAGCT
 GTAGGTTTGAACCTTTGGAGGTGGTGGGTGGGGCTTTGTGATGGGATACC
 CACTCATATCG

FEUILLE DE REMPLACEMENT (REGLE 25) CCTCCC

FIGURE 10

SUITE 31

204770-097E402

ACCCTCTTTTAGTTCCTCTTCTTGGTTCTACAAACACCAACCTGTGTGTA
 TTTTGGTGCTGCCTGTTCTCTTTTGGGCTTTCTCAGAAGAAAATGGGTT
 TTTGAGGGAATCCATTTCAGGTGAGTCCTCACCCTAAGCAGCTCTTCTTCA
 CTTTGTGGGCCCCAAGCTGACCCAGAGCCATACACCCAAAGCAAACCCAG
 AGCCGTACACCCATAATGAGGCAGGAAGTGGAGTGTGCAGAGCACATCTT
 TTAATTAATAATTAATATCAGAAACGTAGGCAGAGACCAGCTCCCCACAC
 CAGGCGTTGCTATTTGCAGTGAAAGGCCGCATACCTTTGCAGGACACCCC
 AGATCTGCCCCACGATTGATGTCAAATAGATGCATAAATTTCTTCCAAG
 TCTTCAGTGCTCTCTGGTGGTTTCCCCACCCTGCAGAGGGACCGCCCCGG
 GGCTCCCCAATGGGGACAGACACAGGGCAGAGCAGCGGGTCCCCTTGGCAC
 ATTGCTCCAAGCAACCACAGCACACATCCCATCAGATGCCCTTTTCATAA
 AGGACATCTCAAGGACAGATCTTTAGGGGAGATCTAAACCCAACCCAATC
 CAAATGGGACATCAGCTGCCCACTCGTGGACTGCTCCTCTGAGGGGGGAT
 TTTGGGTGATCTCTTGAAGCGAGCCCCCAGCCCTATCTTGAACAAGGGG
 AGGACCTTCTCCCCATTGAACAAAGCCCTGGTGTACACCAAGATGGGGGT
 GTCATCATCCGAGCTGAAGAATGCCACCCGACCCCTTCGTAGTCCAGGG
 AGACCCGAATCCTCCTGGGAAGTGCATTACAGCGTAGGTGGCACGGGGGA
 GACGTGAGGGAGTGGTAGGCCCTCCAGCGCCAGACACCCTCTTGGGGGT
 GAAGCTCATGGGTCCCTTCTCTCATCGAAGCCCGGGCCACCCCGAGG
 CCCACACCCCCCTGTCCACCTCCACCTCCCAGAAATGCCTCCCCGAG
 GTGAAGCCCTGGCAGCCCAACACGCAGGGCTCGAAGCTGAACCTCTCGGG
 GTTCTCGGGGAGGTCTGTGGCACCAGTTGGCCCCGGGCTTGTTCGGT
 CTTCAGAGAGATGGAGGTGGGGTGAGCGGTGGTGGGGTCCATGGTGACG
 TTGGCTGTGGGACATGAGGGGGAATGGAGGTAGGATTTAGGCTTGGGGG
 AGCTGGAGAGGTTCTCTTCTTCTGTCTTTCTCTGGGTGCTTTTGA
 CATGGGCTGGTGGTGGTGGTGGTGGTGGTGGTGGTGGTGGTGGTGGTGG
 GGTCTTTTCCAACCTTGTGATTCTATGGGGTGTGTGGGGCTCCACCAGC
 CTCAGTGTCCCCAGTAGAGATGTAGGAGAATGGGGAGAGGACAAATTTT
 AGGGCAGCATAATGCGGGAGGGACAAAGACATGGGAAGGGGACAGCTTGA
 CATTACAGGAGGGGAAGGGGAAGCACAACACTGTTAGGTTTTGCCTTGA
 ATCTGTTACTGGCTTTGTAGGACCACCAGCATCAGGATGCTGTCCCCATT
 CCTCCCTTCCCTGTGGGACTGCGTTGTTTTTTCCCAAGAAAACCACTCC
 CCACCCACATCCACCACTGCTGACATACCTGGCTCTTGCAATTGAAACA
 TCAGGCTGTCTGAAAAGGAGAACAATTCACGTGCAATTGGGTTTATGCTTC
 AGGAAAAGGGGCTGGGAGATGGGGAAGGGAAACCATGGGGGTCTGGGGG
 TTCGAGTGCAAAAGCTCTGGGTTTACTGCAAGAGCCCCACGACCCTCCC
 AGACCTGGAGGAGACCCCGACCCCATTCAGTACCTTGGCACTTCTGCAGC
 GTCAGTCTCACCAGGACGTTCTTCTGAAGGAAGTCCTCCAACCTTCTTTC
 CAGAGTGGGGGAAATCTCTGCTGGAGGGCTGAACTTCATCATCTCACAGC
 TGCAAAGAGAGGAGAAGGGTGGGGATGGGGGGACTGTTGCCTTGGTTGGT
 TGGCTGTTCAATTTTATTCTCAATAGGAGAAGCTATGGGGTGAGGATATTT
 GCACAGGGACGAAATCCCTTTCCCCCTGGGATCCCTCTGCCTTGACGCC
 CTCCCCAGGGTGCCATCCAAAATCAGGGTGACAATAGGAAGGAGCCAT
 GTTACCTATTCAAGAGCCTCCTGATGTCCTAAAGGTGGGAGGAGAGAGGA
 GAGATGGATCAGAAGAGGAGCACCAAGGGCTGCCCTTCGTATGGCAATG
 CACAGCAAAGACCACCCTGCCACGGTGTGATCCCCCCCAGCAGCAACAC
 AGGGAGCTCCCATGGGGTTGAGTTTGGGTTCTCAGGGTTTGCTCTGTCCC
 CCCATTTCCACCACCCCTTTGGGTTCTCACCAGCAGGAATTTGCTGTGC
 GGCTGTGGAATTTGCCCTCCATCTCCAGATCAGGGTGTCAAGGTGGGA
 CATCTCTCTCCATCACCTTCGTACCGCATCCTCTGTACTTTGGTGACGG
 CTCTGTCCAGGTCTGCCAGCTGGACCAGCAGGAAGCGCTCCTTCTCCTTC
 AGAAATCGCTGCAACTGCTCGAATTCACACACTATCCTCTTCCCTTCCTT
 CTTGGTTTTCTCCTGTTGGGATGAGGGAGAAAGCCAATGGGGTGGAAATAG
 AGGCAGGAAGAACCCCTGGGCTCTCAGGATGCCCTTTCTGGGGGATA
 TCCAAACCAAAACCAATGGGATGTACACCAATGCCATGGGAGCACAAAC

FIGURE 10

SUITE 32

ACTAATGCCAATGGGAATTTATCACCAGTGCCAATGGGAACGTAACAACA
 GCGCCAATGGGAACGTAACACCAGTGCCAGTGGGAATTTATCACCAGTGC
 CAATGGGAACTTAACATCAAAAAGCCAAAGATCATCTTGCTGGGCATTTG
 GGAGCAGCAGGAATTTTTCAGGAGTTTATCCCAAAAAGCAAAACCAAAGG
 AGGGGGTAGGAGATGAGCTCTGTATGAGGGATATTTACAGAGTTTAGGAG
 GATCTGCTACGTTATCTCTTTAACACAGGGGTTCTGCGTAACCCAGCT
 GATAAACACAGCCTTAGCGCTTTCCAGCCAGCTGCGAGCCAAAATGC
 ATGATCTGCCCCAAAATACACCAAAAACAACAGGACAGGGCGGAGGGGA
 AGGCAGACACCTCCCCTGCTGCACCCACCAAAATACAAGCCCGTCCTTCCA
 CCAGTCCTTCTGCTTTTCCAGGTACTTTTCCCTCTCCTCCTTTGAAGCCTG
 GAGGCGAGCCTGAATTTCTTCTGTGCCAAAAGAAGAAAGGCGGAAAGCC
 TGTTTTCCCACTTAAACTGCTTCTGTGAGATGGGAGAGGCTTTGCTAAAG
 CCTGGAATCCTCTGCAAGGTGCAGAGCTGGGCAGAGGGAAGCTCTGTGAG
 CACGGTGTGCTGCTCTGGAGCTCTGTGCAAGCTGGGAGTATTTTGCAGAG
 AGAAAAGAGGGGAGAAGGGAAGGAAAAACACGAACTTGCTGCAAACGTAG
 AGAAAAACGCTGCAAAAGAGCAACAAAAAATCAGCACTGACAGCTGCGC
 AAGGAGGTGTGGAAGGGCAAGATAAGCACTTGGTGAGATTTCCCTCATAA
 ACACCCAAAACGGCGGCCCTGGGGTGTGTTTCTGTATTTAAGAGCCCTC
 AGTGGAATGGTTTTTGCAGGGCTGTGGTGAAGAGCAAAGCATCAAAGGA
 AGGAGAGGGCAGTAATGTTGCAAAGGGCTGACGGCGGTGGTTGCAAAGAG
 GGAGGATGGGGGGGATGCGCCAAGCAAAAGGTTGCGTGGGTTTACCCGC
 AGGGATGCACTGCGCCCTTGGCTCCGGGTTTTGGGACCGTACCTTGTA
 CCTGGGCCGCTGGTGGGCAGGGAGCACAGCGTGGGAGCGGTGCGCCTGG
 GACGCGTCGCACTGCGCGCAGATAGGCTCTTGGTCTCTGTGCAGAAGAG
 CTTGAGAGCCTCGCGGTGCTGCTTGACCAACCCGAGGAATGCAAACCTCA
 GCTGCCGGGCGATGCTGGCGATATTTGCCAGCTCTCTGCTGGGGCGGAAA
 TTTTTGTGCAACGCCGTTCCTGCACTGCGGACAGGGGAAATTTCCCTC
 CAGCCCTTCCAGCAGCGGGCGATGCACTCCCGGCAGAAAGTTGTGGCCGC
 AGGGGATGGAGACGGGATCCTGGAAGTAACCCAGGCAGATGGAGCAGGAG
 GCTTCGCTCTGCAGGCTGTCCAAGGGGCTCTGCGTGGCCATGGGCTTCCT
 GCTGCGCTCCGATCCGACAGGGAATAGGGGACCTTTCTCCTTATCTCC
 TCGCTGATAGGAGAAATCCGGCCCCGGAGGCTGAGCCTGAGCCAAACAGG
 GCTGGGAGAGCTCAGCCCATAGGGGATGCTGGTGGGAATGGGGGCAGCTC
 GCGGCTCCCCAGCACGGAGTCACCAAACCTGGGGGGATCTGGGGGAAATTC
 GGAGGAAAAGTCAGATTTTGTCTCTCCTCGAGCAGCAAAGAGGGCAGGG
 GAGGCGATTTTCCCTTCTGTGCGATCACTGTAAGGAATTTCCAAAGAAA
 ACGCATGGAGGTCTGCTTGTGGGATGGAATATAGACGTATATTGGAATA
 AATACAGGAAGACGTTGGAACATGGGAAGGCACTGAGATATAAGCGTGCT
 GTGTTGGATATGACTCTGCTCGACTAAAGTGAAGGTGGTTTTAATAGCAC
 TGCTCAGAGCCAGGCGGGTTTTGGTGTGTTTGGGGGGAATTACGTGGGT
 TTGGAATTGGGAAATATGAGACGGAAAAATAAGAATAATGGAAGCGCCCA
 ACGTGGGGCTCGAACCACGACCCTGAGATTAAGAGTCTCATGCTCTACC
 GACTGAGCTAGCCGGGCTGATGGGCACGCACCCTTCTAAGCAATACTTCA
 TGGTGATCCTGCGGAGGGGTGCTAATAATTCTACCTAATTATTTTGTAA
 TTATCCCGGTAATTATGGGTTCTGAGCAATCGGAATCCACGGGGAAGAG
 CTGCATGGGGAAAAAGCACCTATCCCTACGGGAATAGCCGGGAACTGCCC
 GGCAGTGCTGCAGGGCGGGGAAAGAGGGGAAAAAGCAGGAAAAAATGGG
 CAAAATGGAACGTTTAAAAGTGGAGAAATTAACAGTGAAAAAAATGCAGG
 AAGCGTAAAAGTAAAGGCTGTGTTTCTGCCCGGTTTCGAACCGGGGACCT
 TTCGCGTGTGAGGCGAACGTGATAACCACTACACTACAGAAACGCGCTGA
 AGGCGGCTTCGCCGCACGGAGATGTGAAGGGGCGAATGCCGGGGCTCGGT
 GCGGAGTTTGCAGATAGGGGCGGCTCCGGGCCCTCCCGCGCGGTTCCG
 GTGAGCAGAGAGTGACGCGGTGACAAAATGAAGGGAAAAATGTAAACT
 GATGCTCCCGAATCGAGGCTCGAACCGCCATTGTCCGACTGACAGCCGCG
 CGCTCTACCGA

FIGURE 10

SUITE 33

FICHE DE REMPLACEMENT (REGLE 125)

CCGTAGAGCGCCACCCCGTTGCCTAGTGACAGGAGCGCCGCTTCCGGTC
 AAGTGATGAGCGGAGGGGGCGTGGCTTGTGTGATAGGACGGAAGTTCC
 GGTGAGGTGGTACTGGAAAGGGGGCGTGGCTTGGCGCAAAGGGGACGGAA
 AGCGGAAGTGCTGCCGTTGGTTGGCGGAGTTCCGACCATAGAAGAACGAC
 GGGCGCGGTGGGAGGGCGGGAGGTAGAGCGGTCCCCGGGGAGAGTGCTGA
 GGGGAGCGGGCAGGCCCGAGGAGGGAGCGGAGCTTACGGGGAGTGCGGAG
 CCTCGAGGCGGGTCCCAGCGCTTCGCTGTGGGGCAGGAGAAAGGCTTCGG
 GGCAGGAGGAAGAGGGCCTCGGGGCCTCCCCATGGAGGCGGTGGGCGACG
 ATGGGGCGTTCGTGGGGCGGGTGAACCCGGTGGAGACGCTGCAGGAGGAG
 GCGATCTGCGCCATCTGCCTGGACTACTTCGTGGAGCGGTGTGATCGG
 CTGCGGGCACAACTTCTGCCGGGTGTGCATCGCGCAGCTGTGGGGTGGAG
 GAGAGGCTGAGGTGGAGGAGAGCGGGCGGGGCCGCGGCGTTGGAGGAGGAA
 GAGGAAGAGCTGGAGGAAGAGGAGGAAGATGAGCTGGGGGAGGAAGAGCT
 GGACCTGGAGCAGGAGGAGGAGGAGGAGGATGGAGGCGGGGAGGAGGAGG
 AGGAGGACGACATGTGGAGCGAGGAGGAAGAGGATGGAGAGCTGTGGGAA
 GGTACTGGGGGTGGGTTTGGGCGTGGCCTGCTTGTGAGTGTCTTTATGGATGA
 GTGAGGGAATTGGGTGACCCCTCAGTCAGTTTGCAGATGATGCTAAGCTG
 GGGGGGTGTACTGATCTGCCTGAGGGTAGGACGGCCCTACGGTGGGGTCT
 GGACTGGGCCCCGATGGGCTGAGGGCAATGGGGTGGAGTTGAGAAGGACCG
 AGTGCCTGGTTCTGCACTGAGGTCAACAACCCCATGCAGCTCTACCTG
 GGGTAGAGCGGCTGAAAGCTGTGTGAGGGAAAAGGATTTGGGGGTGAATA
 TGAGCCAGCAAGAGGCCAAGAAGGCCCATGGCATCCTGGCTTGTATCAGA
 AATAGAGCAGCTAGTGGGAGCAGGAAGTGAAGTGTCACTCTGTACTGGCAC
 ACCTCAATGCTGCACCCAGTTCTGGGTCCCCTCTCACTACAAGAAAGACA
 TTGAGGCCCCAGTGAGGATGGTGGGGGTGGACTCAATGATCCCTGAGGTT
 TTTTCCAACCTTGATGATTCTGTGATTCTCAGACCCCGTGGAAAGAGGAGC
 TGTGGGATGGAGTGGTGCAGGGAGAACTCTACTTTGGGGACGATGATTAT
 GATGAGGATGTGATGGAGGAGGATGTGGAGGAAGAGGAGGAGGAGGAGGA
 TGAGCGCAGAGCCCTCCGCCCCCTGTCTGCTGCCCCGCCCTCGCCGCC
 TGCAGACCTTACCTGCCCCCAGTGCCGCAAAACCTTTTCCAGAGGAAT
 TTCAGACCCCAACCTCCAGTTGGCAAACATGGTGCAGATCATCCGGCAGCT
 CCACCCGCACCCGCAGCGCCTCGCGCCGCCCGCGGGCCCCCTCAGCCTCAG
 GGGGTCTTGGGGGGAACCCAGGGATCCTGGTGGCAACAGGAGGTGCGGGG
 TGTCCGAATCTGTGCGAGAAGCACCAGGAACCCCTGAAGCTGTTCTGTGA
 GGTGGATGAGCAGGCGATCTGCGTGGTGTGCAGGGAGTCACGGAGCCACA
 AGCATCACAGTGTGTGCCCTGGAGGAAGTCGTGCAGGATTATAAGGTG
 GAGTTTGGGGAAGGGTCACGGTGGGATAGTGGGTGAGGTGGGGTTTGGGG
 AAGGGCTGTGGTGGAGAAGGCGGGGTTTGGAGGAAGAGTTATGGGAGAGT
 GGAGGCTTGAAGGGAAAGTGAAGTTGGGATCAAGCTAGGTTCTGTCTTGCT
 GAGCTGGTTGGGTTGGAGGCGTGGGAGGCTGGGAAACCACACACTGCAAT
 GAGGAGGTGGAAGGTTCTGGGTACCCATTTTCTGCTTAAAAACACCTTCC
 CAGCACAGTTCCTCAGAGAAAGCAAAGGGAAGTGGCGTGAAAGTTGGCT
 CTGAGGTTCCGTTTTTCACTCTGCCACCAAATTAGGGACAAAAAGAGGCG
 ATGACAGAGGGGATTGCCCCAGGCAGGGTTTGTGCTGAGTTGTGTTTCCCTC
 CCTCAGTACAAACTCCAGAGCCATTTGGAGCCACTGAAGAAGAAGCTGGA
 CGCGGTGCTGAAGCAGAAGTGAATGAGCAGGAGAAGATCACAGAGCTGA
 GGGTAAGAGCTGAAGGTTTCTGTGCTTCATAGAATCATACAGGAGAACCA
 TCAGGGTTGGAAGAGACCACAAAGATCATCAGTTCCAACCATCACCGCTG
 CTGGGAGTGTGCCTTGGTGGCTGAGCAAGGAGAGAGAAGCTTTGCTGCTG
 CTCTGAGCTCTCACGGAGGCATCATATTCCTTTCTGCAATTATTGGGC
 TGTGAGGGCTTGGAAACGGTTTCCAGTTGAATTAGAGCTTAATGAGAGC
 TTTGTGTGCCTCAGTGTTGAGTGGGAATTGGTGGTTTGGGAGCTGGTATT
 CCTCATTGTAGTTGAGGATGCTCTACATCTCTAAACCTGTGCAGACTTTG
 CTCAGTTCTGTCTGTGGTGCATTGAGGAGATGGTAAGCTTATGGTGTGT
 GGTGAACCTGAGAGAAGCATAGCACAGCAGCCCCAAATGAGCTGATCTC

FIGURE 10

SUIITE 34

10043160-034402

TCACCTCCCCCTTCTGCAGCAATTCCCCTAATGCTTTTCCCTCCCTCTGCA
GGAAAAGATGAAGCTGGAAATCAAGGAATTTGAGTCTGATTTTGAGCTGC
TCCACCAGTTCTCTCATTGGGGAGCACGTGCTGCTGCTGCACCAGCTGGAG
GAGCGCTACGAGAGCCTGCTGGCCCCGGCAGAGCAGCAACATCAGCCAGCT
GGAGGAGCAGAGTGCAGCCCTTAGCCGCCCTTATCACGGAGGCAGAAGATA
AGAGCAAGCAGGACGGGCTACAGCTGCTCAAGGTCTTCTTCCATCCCTTT
CCTTGTCTTTATGGCAAAGCGATAGCACGATGGTGGGAATAATGCTCCAG
AAAGCTTCTGTGTCTATGAGAGAGTGCCTTTAGTTGGTGGGCTGGGTGCTT
CTCCACCCCTCCTTGTGGTGGTTTTTGGGGAAAATGCCGGGGGGGGGGGG
GGGGGGGGGATATGCCCTGAGAGATTTAGGGTCTGTTTTGGTAAGGAAAG
CCTCCAGCAATGTGTGGGCTGTGTCTTTGTTCTCTGTGGGGAAGGGAATC
ATCCAGGCTCAGTGTGAGTTGTGGCTGATAAGAGGATTTATTGGGAGCA
ACGGTGGGATTGGTATCAGTCATCCCTAATCCTTTCCCTTCTCTTTCCCAG
CTTGCTGCCCTCCTTCCACAGGACATCAAGGGCACTTTTATCAGGTGAGT
GACTTTGTTTGCATCTTTTCACTTTGAATAACTTTTCTTTTTTTTAAATGT
CAAAAAAGCATTGAGCTTTTGTTTTAAATCCTGTGTGATGGGTACAGTT
GGGGCCTGGTAATGCAGGGGAAAGCTGTGTCTTAATTTTGGGTGATGGA
AACTTCTGGCTGATGGGGTGAAATGGGATCTGGGGAACAATTTGGGAAA
AGACTTGGGAACTTGGGAAACAATCTGGGGCCATTTGGGAAAGGGGAAG
GGTGGGGAGGAGATCTCGGCCCTGATTTCTGGAAGCGTGGGTGTGCCCAT
GCAGACCTCATGCTATAGCGAAACTCCTCACTCTGGAGAAACGATTCTCC
CCATCCTGTGAGACAAATGGGCAGCGCTGGGAGTTCTCAGCCATGCTGGA
CGCACGTGGCTCTACCCCACTCTGTCTGCTGGCTGAGGGAGGGGTGGGGG
AGGCTGGCTGCACCAGTGCACCAGTTTGGCCGATCCATGCGTTGCTCTG
GTTTTTCCAGAGCTGCATGCAGGCCGCTCACTTCTTTTCTGCTGCTGAA
ATTCTCTGCTTTCTCCTTTCCCCCACCACAAAAAGATGTGAGAACATC
AAATTCAGGAGCCCCGAGATGGTGCTGGTGGACGTGGGGAAGAAATACCG
CAACTATTCTCTGCAGGATGTGGTGATGAGAAAGATGGAGAAAGCCTTCA
GCAAAGTTCCACAGGGGTGAGAGAGTCTCTCTCTCTACGTGGGATGGGG
TTCCCTCCACTTGGGATGGATTCTCCAGCTCTCTTGGGGTTCTCCTTC
CATCTCTGTGCTCCCATGGTTTGCAGCCTGATGATCCTTTAGGAAAAGCA
GCATCCCTCTGTTCTCTCTGTGCTTTTCCCTTTTGCCCTGTCTGGGTTT
TCCCTATTGTAGCTCCTCCATAGAACTGGGGTTGATGTGGATCTGGATT
CATTATAAAGGAGGGATGACTGCCTCAAACCTCAGCATGGTGCAGATACGC
AACCAGATGAGGATTTAGGACTGGGGTGCAAGGGGGGAAAAAGTGCCAGG
TGACCCCTAACGACCCCGCTCTCTGCCCTTCTTCCAGCTGACATCAC
GCTGGACCCGGACACCGCTCACCTCGCCTCAGCCTCTCCCTGGACCGCC
GCAGCGTTAAGCTGGGAGAACGACGCCAGGAGCTCCCCAACACCCCAA
CGCTTCGACTCCGATTACTGCGTCCTGGGCTCCAGGGTTTCACCACAGG
CCGTCACTACTGGGAGGTAGAAGTCGGGGGCAAGAAAGGTTGGGCGGTGG
GGGCTGCACGCGAGACGGCTCGACGCAAGAAAAAACCATGGGGCCTCAT
CAAAAAAGGGAGATCTGGTGTGTTGGCACCAATGGGAAGAAGTACCAAGC
GCTGACGGCCATGGAGCAGATGGCTTTGTCACCAGCGAGCGGCCCGGC
GCTTCGGTGTCTACCTGGACTATGAACGGGGTCAGCTTTGCTTCTACAAC
GCTGAGAGCATGACCCACATCCACACCTTCAACGCTTCTTCCACGAGCG
CATCTTCCCTTTTTCGAATCCTGGCTAAGGGCACTCGTATCAAAATCT
GCACCTGATGGCCCTCCAGCTTCTGATTTTTTTTTTCCCTTTTTTCCCCC
TGCCTCATCCTTTGGGTCCCACTTTGGGACCAGACGCTGCATTGTTGTC
TCGCACCTGCTTGTCTACAAGGCCTCTTCCCTCCTCTCTCTGTCCCAGC
CTCTGTCCACGTCCCAACTCTTCTCCGGGGTCGCGATCCAGGCTGGTTT
GGTTTGGAGAAGGGATCCAATCTCCTTGCTGGAGGTTTCCCTTCAGCTC
TTGGTGTCTATGGGCTCCCTCTGCCTTTCCCACTCCTCGCAGCAGCTTTC
CAGTGTGCTCTTCCCGTTTTGTTTAAAGCCTGTGGTGGAGCTTTGCGTT
GTTTGGCCTCTTTGGATGCAGAGCTCGAGCTGAGGATGCTGGGGTCTGTA
CATTGTGACACGAGCACTGCTTGTGCCCTCTTGGCCATTGCTTTCTGAAA

FIGURE 10
SUITE 35

204779" 084407

2043160-01402

GTCACCTCAGATGCACCAAGGAGCCTCATTTCTTTTTATTTTTCAGTTCTG
GGGCACAACCTCTGCCCACCTCCCACCCAGCCACCATCTGGACCTCAAA
CCTTCCACGTTCTCCTATTCTGCCACTTGTCCACCTTCCCCTTTTGCTCT
TCTTCCCCCTCTGGGGGTCTCCAGCTCTCCCTCTGCCCCATCATTCCCTC
GCCAACCATTCTTGTGGGCCTGGCACTTTATTTAGGGGCCACGTAGGCCG
GGGAGGGTGCAAAAAATTGGGCAACTTCCACCTCTGAGGCTGCTCAGAGT
GCAGCATCGCACCAGGCCGACCCGGTGGGAAGCAGCCTTGTTTCCCCTTG
CAGCTTAAGAGCTCTCTGAGGTGGGGGTATTTATTTTCTCTTCCCCTTTC
TCAGCTGCTGTTGAATTTCCAGCTGAATCCTGTCCCACCAGAGAGACTCT
GATTGCACCCTGTTGTGTTTTACTTCTTTTTGTTGGTGGATTGGTATTTT
TTTTTCTGTTGGCGTTACAGAGCTAGTTCAAAATATTTTGGCTAAAT
AAGAATTAATGGAGATCTAGTTTTTGAATGTCAAGAAATAATAATAA
TAATAATAAAGAATAAAGAATAAAGTTTTAAAGCTGAGCCTCTCCCTTAT
TGAGAGCCCCCAGGGGACAGGAGTTGTGGTGCAGGCCCCCAGTCTGCTG
TTAACTCCTGCTGGTAAGATGTGACTTAAGCCTTGCAATCGTTAATCTTAA
CTTAATTAGCAGTAATTTGGATTGGGCTGCTTCCCTTCAGCAGCTTGTA
AGGGATAGAGGCTGCTGGGTGAAGTGAAGTCTGTGTTACCACCTCTCCTG
CTCTCCCCACATGTTTTTGGTGGTGGTGGTGGTCTTTTTTGGCCACGGC
TCTATCTCCCCAGGTGTGCACTCACTGTGGGCTGCTACTGCTCCTGAAAG
GGCTCAGGGAGACATTTGAGTCCCTTCGTCCACACGTGGGAGGAGAGCAC
TGATGTCCCCATCCTTAAAGTTGTGGGCACAGCCTTGGTGGCAAATCCAG
AATGGGATATAATGCAGCCATGAGCTCAACAGAGCGCTCTTTTATTGAGT
TTTGTGCATAAAATCTGTGTGTTGTTACCACATCCTCATCTGGTTCCAAT
GGTGAAGTGGCACACCCGGACGAGGTTATCTGTGTAGCCAGCAAACAGCG
TCTGGGGAGAGAAATGGAGGAAGTGGATCATGAAAAGATAGGAATCAGCC
CTCGGTGTGAACGTAAAAATCTCAGAAGGCAGCTCCCAAAGCGGAGGTGC
TGGAGGAAGGTGGGAGTTTTAAGGCTGCAGGAGGAGCAGTAAAAGGGAA
AGGAGAAGGGGATATTTCTACCTGCCCCATCTGCAGACCACGCCAGAGAGG
TACACTGGGGAGGCTCAGCTTTGCTGCTGGTGTGATCACCTCCTGCTTC
AGCTCATCCACAATGATTTTGCCTTCCAGGTCTGTGCAGGACAGAAGAG
AGCGTGAGGGACTAAGGTCTGCAGGGAGACTGCTGTAGCCAAACCCAAAC
CATTCAACTCAGAACAGGCTCAGGGTGTCTCAGAAACAGCCTCTGGGTTT
CCGCACAGGGATGCAGTCAGATGGCATCGAAGTTTCATCACAGCAGAGTG
GTGGCTGTGCCCCACACCACCTCCAGTCCAGGGGATGACAGTGCCACC
AGCATGACCCATCCCACGTAACCAAAAGGGCTCTGCACCAAGGCATCTGT
GGGGCAGGGCGAGGATTTGACCACAACTCTGCCTCCAAACCCACAGGAT
AAGGGAAGTGATTCTTTAGGAGGTAAATAGGGATGTACATACCCAGATC
TTGATGCTGGGGCCGGTGGCAGCGCAGAGCCAGTAGCGGTGGGGCTGAA
GCACAGCGCATTGATGATGTCCCTCCATCCAGCGTGTACAGGTGCTTGC
CTTCATTAGGTCCCACAGCATGGCCTGGCCGTCTGGGGGGCAGCAAAG
AGGAATCACAGCAAACCATCAAACCTGTGGCTTTGTTCCAGTTGTCCATC
TAAAACCTTCCAGCTTGGAACAGCACTTGATTTGTGACTGAGATGTGGG
TGAGTTGCCACAGGACAGCAAGAGGCACATAACTGAGCTGTGAGAACAAC
AGAATAAGCTGCAATTTGGCCTCAGCTTTCCCCCAGGGTGTACCTTGCT
CCAGAAGCACAGAGGGAGCCATCAGGGGAGACAGTCACTGTGTTTCAATA
TCCCGTGTGGCCGATGTGGTTTGTCTTCAGTTTGCAGTTAGCCAAGTTCC
AAACCTAAATGAGGGTAAACGTGACAGGCTCAGAAATATGGAGGAGAAAA
AAAACAACCTCTCATGATCACTGTCAAATATTCCCCAGAACGCCGCAC
AAACCCCAAAGGAGCTGCTCCTCTCACCTTCACAGCTTGTCCCAGCCAC
AGGAGACAATGATGGGGTTGCTGCTGTTGGGGGAGAAGCGCACACAGGAA
ACCCACTCAGAGTGGCTCTCGTCTGAGGAGAGGAACAGCATTGGGTTGA
AAGCAATGAAAAGCATCCCCAGTCCGAGCTGCTGCATCCCACTGCTCCCT
GAGCCCCCTCATAATTGCAGGACGTGTCTCAGACCCCCCCCCAGAAAGAAA
GGTCAGCAGGCACTGTGTCACTTCTAATCATTAGGACGGAGCTGGGAGAT
GTGGATTACCGAT

FIGURE 10

SUITE 36

ACTCAGAAACAAGCAGAAAGAGGTTTATTTTCACAGTGTGGAAACTCAGA
TCCGTTGCCTCACCTGCACCGTGTATTTGCAGACACCCAAAGTGTCCAG
AGTTTGTATGGTTTTGTCCCTGGAGCCCGAAACGATCTGGCGGTGTCCGA
GGAGAAGGCGACGCTCAGCACATCCTTGGTGTGGCCAACAAAGCGGCGGG
TGGTGGTTCTCTGCAGGGACACCAGGAGGTCGCACGGGAGGGACAAAG
CTCAGCAAACCCCATTAATAATTAACCCTCCCCTAAATTGAGGAGAT
CGTGCTGCAGTGCATAAATCTTAATGAACACAACCTGATGGAAGCAGGAA
GGAAGCTAAAACGGAGTCATCTCCACATGGGTTGAGGAGTGGTGGTTCCCT
TCCCTCCTTCCGAACAGGAACAAAAGGGTGCCAAAGCTTTTGATATAGGG
TTGGAATAATCATGAGGAGTTTAGGATATAAAACTCAGCTTCCGTGGACA
CACAGCAGCGTAAGTGCTGAACGCTTTTGGAGGATTGGGGTAGTTCTGCT
TCTTGAGGAGTTTCTTCTCTATAGTACTCCCAAAATCAGAGTGCAAGA
AGAGCCGGTGTCTCCAACCTCACCCCAAACCTCTGTACCCCAAATCAC
ACCGAAGGAAAAGCCTGCTTGCTCCAGTCTGTACCCACAGCGATGGTGA
AGGAAGAACCAAATCCCCCCTGCTGCTCCACCTGCTTCTCTCCCATCAT
AATTGCAGGACGTGTCTCAGATCCCGGAGGATCAGCAGACTGTGTCAGG
TGTAATCACTGGGAGAGTGAGCTGAGGGAGGAACCGCTTTGGTCTCCCT
CCAAGCATGATTTACCACCCAACCTGAGAGGAACTCACCTCATTTTCAGC
CTGTACCGCACACCTCTCACCCACCCCAACACCCAAACAAAACACAGAGC
CCAGCTCTGCCCCAAACCCCCACCCCAAAGCCCTTTCAGTCCCCAGGACT
CACGTGGTGAGGTCCACAGCCTCAAGGTGCCATCCAGGAGCCCGACAG
CGCAAACCTGCCCATCGGAGGAGATGACCACATCGCTGACAAAGTGCGAGT
GGCCGCGCAGGGCGCGCTGCGGGATCCCGTAGTTGGTCTCATCTCGGGTC
AGCTTCCACATGATGATGGTTTTGTCTGGGAAGGGGAAAGGCAGCGGCC
TCAGCTCCAACCCCTTCTCACATTCCCGTCTCACTGGGCTTTATCTCCCT
CATAGCAATGGGGGGGTTACACAGAAGCACCGCACCCCTTCTCTCAGCC
CCCCAACCGCCTCCCTACGTCTCTACACAGCAGCCTCCCCACCCTGCA
GCTCTCTGTCCCCGAGCCCTGCACCCCATTCATCACCTCCCCTCCCCAT
GGTCCCCCCCAGCCCCCTCCTTACCAGTACGGTCTCCCCTTATCTCCC
ACAGTCCCCTCCATAGGCCCCACAGTTCCCTGCCCCCCCCACCCACAG
TTCCGCCCCCCCCGCTCGGACGAGGCCCGAACCCTCAGGCGCGGCCCT
CACCCCGCGACCGGAGAGAATCATGTCCGGGAACCTGCGGGGTGGTGGCG
ATCTGCGTCAACCCACCATTTGTGGCCCTTCAAGGTACCGCGGAGGGTCAT
CTGCTCCGTATGGCGGCGGCGGGGCGGAGGGATGGCGGCGGATTCAATA
AAGGGCCCGGCCCGGTCCGGTCTACCGCCCGCGATGGCCGCCAGCGCGG
AAAGAGAAAGAGGGAGGTGACTTCCGGCGGAAGCGGAAGTAGCCGCTGGG
TTGTACGGCAAGAGGGGCAACATGGCGGCGCGCATAGAGAGCACGCTGAA
TGGGGGAATGGGCTTTGGAGGTGGGGAGGGAAGGTTGTTCTCTGCCGCTG
CAGGGACACGAGGTGCGGGCAGAGCACCTTCTTTAACATTTGCTATTATT
TAACGTTTTACATTTAGCATTTTTATTATCCCTGTTGTGCCAGGACGGAG
AAGAGCAGGGTGTGCAGCCTGTGCTTATCACCTGCAGCTGTCCCTGCACC
CCACAGCCAACCCAAGTTTGTGACGCCTGAGCAGGATCTGACCCAGGAAG
GCAAACAGAAGGTCTGAGTCTCTCCCTTTCCCTTCCCATCCCTCCCAC
GCTGCAGTTTGGGGGCTGTGACCCGTCCGCGTTGCTCAGTGCTCATTCGG
ATGAGCAGTGGCTGATGGTGATGTTACAAAGTTTTTGGCATCCCTGTGGG
TTCCACCCCGTTTTGTCTCACCAGCCTTTTTCTATCCGTCCCTTATCAGC
AGATCATCCTTGTTATTAGATCTGTCTTTTTCCAGTCACGGCTTTGCATT
TTCACCTTGGTTTTACCACCTAACATCAAGCCTTTTGTCCCCATCTGATG
ATATTCTATGCAGATAAATCCGTAAAGCAGGGAAGATTAAATTCTGGCCC
CTTCTACACCCATTAGGTTTAGATCTTTGCAGCATTCAGCCAAGACGTG
CTTCCAGAGCCAGGAATAACGTGTCTTGATGTGCCAACACACCTTGAAT
CCAGAAAATTGCCCCAAAATAGGCATGACTCAGCAAGCACCGTAGTGGGC
ATGATTTGCTTGGGTGACCCCGTGGGTAAGGAGCCATTTGTTGGACACCA
CGATGTGCTTTTTACAGCCCTGTGAGCGCAGCGTCTTAAATTGCCCTCC
AGACATTCCAAATTTGGGAGGCTGAGGAGGCTTAAAGGAGGCGTCA

FIGURE 10

SUIE 37

GCCCTCCGGGATGAAGGAATCTCTGCCGGGGTTTTCCGTTGGATCACAGC
 AGGAGGATTTGCTTTCTAAAGCATTAGAGTGACGTGGAGAGCCCAAATC
 GGACCCAGTGGCCACATTCTCCCAAGGGAAAACCTTCGGGTGCCCTAC
 GGTTCCTTTCTAGCATGATAACAACTTCTTTTCCATCCGCCCATCCCC
 TTTTGGGTTTGGAGGTTGACAAATCCCACTGAAATTCCTATGTTGCACA
 CATGTCCTTCATTCTTTAAGTAGGAGTTAGCAAAGGTTCCGCATTGACTT
 AATTCAGAGCGAGATCAACAATTTTAGGCATTCTTTATGAACTTCACATT
 GTTTTATGCTGATCAGCAGCAAAAAACATACAGGAATAGGAGTGTGTCT
 GTAGGAGTGCTCTGCATTTTCTTGCTCGTTTGGCTGATTAAGGAAGCTGG
 GAGGAAATGTTGTGAAATAATCCCAAGTGATGAGAGACTGTGGGTATGGG
 AGGAGATGCCCTCTGTCTGGTGAGCAGTAGGGACAGAAGACCTGAGCTC
 ATTTTCATATATCTGTATATTAAGGCAATGCTAACCAGTGCTGTCTGTGTA
 TTTGGGGCCAGGAGTGGCTTCTGCCCCGTTGGTGCCCATAAACCAGTGCT
 GCCCCATTTGGGATTGGGGTCTGCTCGCAGACCACATCCACCAACCAACC
 CATGGCTGATAGCAGAGAGGGCGACCAGGTCAACCCTCCATATATCTCTGC
 AGAAACCTGTTCTGTCTATACAGGGATCCCCATCCCTCCCCCAGCCCTC
 CTTCCATCCTCGGCATTTGGGTTGGCTATAATTAGGCTCTGGGAACGTTT
 CCCTGCTGCCAGCACAGCTGTCTGTCTGCAATGATCCTTCCAGCTCTCT
 GCGGACACGCAAAACCTCCAGCAATCCTAAATACCCATTTCTGCACTCC
 TGGGACAAACTGGGAGCTGCCAAAAATCTCCAGCCCCCACAGACGTGAC
 CATCACAGCACCAAGGAGCAGAGCAAGCGCAACGTGATTACGGTGCAGGT
 CGGGGTAAAGCCTTTCTCTTTCTTCCACAGCCCAGGATTTGGGGGATCCT
 ATTGGCTCTATGGGATCTGGGAGATGCAGGAGAAATGTGATCCCTTTGCT
 GTAGCAAAACAACCTTTTAGAGTCCTGCACCTGAATCTGGCAGTACTGGA
 AAGCAGGAGAGGGATTAAGAGTCCTTCTGCATTATCCTGCTCATAGGGAA
 ATACAGCACAGAAATCATTGGGGCTGCTTCTTTGCTTTCTTGGCACAAA
 TTTAGGTCCTCATTACAGCGTTTCTTTGACTGAGACCCCAATAGGATCTA
 CAGGGGTAGAACAAGCAGACAAAAAGTGATTGATGTTTCTATGCGATT
 TGTTGCCCTTTTCCATTGAGATTTCTGCTTTTCTATGGGGCTTTTTGCT
 TTTTCACAGCTTTTTTTTATTCACTGTAGTGAATAGAAATTTTTAGGGCTT
 TTAGGTCATTGATGCTGTTATGAACACAGAGATGAACTCATAACACCTTC
 CTGGTGTGTTTGTCTATGGGATAGAAAGGAGCTCATGGTGTGTGGACA
 ACTAACAGAGGTGCCTGAGGGCTGGGCCCTCTTTGTGCCCCCTTCTGGGGG
 TCAGCAAACTCCTTTTATTAGATATAAATCCCTCATCCACAATTCAC
 CAGTCTTCCCAATGCAGACCCCAAAAAACATCCCCAATGACAAAGTCCAC
 GAACTGAGAAAAGCAGCAAAAAGCCTCCCAGCCCCAAATATTTATCCCTT
 ATCCCATTTATTTCTATGGGCAAGCTATTCTAGGCATCAGGAAGGTGGG
 AGATTCCAGGTGATTTTGTTCCTAATTGTGATCTTTTAATGATGTTTCTC
 CCATCAGGTGGACATTTGGAAGTGGTCTGACTGGGAAGAGGACGTGATG
 ATGGCATCAGGTAGAGCTCAGAAAGTGGTATTTATCAGCAAAGCAATTTT
 CCAGGTCTGTTTTTTCCCATTTTTTCCCATATTTTTTTCTTATTCAGGGAA
 GAGGAACGCGGATCTTGGTGAGTGATTTTCTTCTTTTACCTTCAAAAAG
 TCCCTTTCCATGTGTAGAAATGGATATACGTACCCCCACTGATACCCAT
 TTCCTTTGTTCTGTCCTTATATTTATACTTCCCCATATTTGAACACATG
 AAAACAAAGCCACATTAAATAAATTATAAACAGTGCAATTTTTGGACT
 ATTATTTTCCATAGAAAAGTATTAAATCAGTGCAGAAAGTGCTCTGGAGG
 TGACTTCTGCAGCACCCAAAGAGAGAGGCGTAGGGCTGAGTGCTCTCTCT
 GTCTCTCTTTTAGAAGAATGGGATGCAAAAATCAGTGAGTGCCCTTTTTT
 CCTCTCCCTTACGGTGAGGTATGGGTGTGGAGGACCTGAATTAATGTGA
 ATTCCTCTGTTTTAAGGGAAGCTAACAGAAGATTTTGGTAAGTCGCTTAT
 TTTCTCGATCTGAGTGCAATTTCTACACCTTTACCATCAGTGATGACC
 AACGTGTGTATGCATTTCTCTTTATTCATTTAGAAGAGAGCGACACAGA
 GCTCGGTGAGTGCTTTGGGGTCTTATCAAGGTGGAAAGATGCCCTCTGT
 GCAACAGTGGGGATTGGGAGAAGCCCTTCAGCTCTTCCATTTATCCACAT
 CTGATACCCAGATGGAGTCAGGATGCAGAACTGGAGGAGGAGGGCCAAAG

FIGURE 10

SUITE 38

1997		1998		1999		2000		2001		2002		2003		2004		2005		2006		2007		2008		2009		2010		2011		2012		2013		2014		2015		2016		2017		2018		2019		2020		2021		2022		2023		2024		2025		2026		2027		2028		2029		2030		2031		2032		2033		2034		2035		2036		2037		2038		2039		2040		2041		2042		2043		2044		2045		2046		2047		2048		2049		2050		2051		2052		2053		2054		2055		2056		2057		2058		2059		2060		2061		2062		2063		2064		2065		2066		2067		2068		2069		2070		2071		2072		2073		2074		2075		2076		2077		2078		2079		2080		2081		2082		2083		2084		2085		2086		2087		2088		2089		2090		2091		2092		2093		2094		2095		2096		2097		2098		2099		2100		2101		2102		2103		2104		2105		2106		2107		2108		2109		2110		2111		2112		2113		2114		2115		2116		2117		2118		2119		2120		2121		2122		2123		2124		2125		2126		2127		2128		2129		2130		2131		2132		2133		2134		2135		2136		2137		2138		2139		2140		2141		2142		2143		2144		2145		2146		2147		2148		2149		2150		2151		2152		2153		2154		2155		2156		2157		2158		2159		2160		2161		2162		2163		2164		2165		2166		2167		2168		2169		2170		2171		2172		2173		2174		2175		2176		2177		2178		2179		2180		2181		2182		2183		2184		2185		2186		2187		2188		2189		2190		2191		2192		2193		2194		2195		2196		2197		2198		2199		2200		2201		2202		2203		2204		2205		2206		2207		2208		2209		2210		2211		2212		2213		2214		2215		2216		2217		2218		2219		2220		2221		2222		2223		2224		2225		2226		2227		2228		2229		2230		2231		2232		2233		2234		2235		2236		2237		2238		2239		2240		2241		2242		2243		2244		2245		2246		2247		2248		2249		2250		2251		2252		2253		2254		2255		2256		2257		2258		2259		2260		2261		2262		2263		2264		2265		2266		2267		2268		2269		2270		2271		2272		2273		2274		2275		2276		2277		2278		2279		2280		2281		2282		2283		2284		2285		2286		2287		2288		2289		2290		2291		2292		2293		2294		2295		2296		2297		2298		2299		2300		2301		2302		2303		2304		2305		2306		2307		2308		2309		2310	
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SUITE 39

TTGCTCTAGGACGGTCTGAAAAGTGACCAAAATCTGCTTTTACTCATT
 TCTTCTTATTTTTTTGTAGCAAAGTGCGATGCAACGATCAGTAAGTGCTG
 CTGCATGTGGGGGTACCTCCATCTTCGGGGTCATTTTCTGCTGTTTCAGCA
 TTGAAAGGACATCAGAATTCCTTAAATCCAACAAAATTTGGGGTCACTCGA
 AGGAATCTTTGCAGATATGGGGGAAATCAGAGCCAAATTTTGAGGGGGGG
 AGGGAAAATCTCAGGGGTGTTTCAGAAATCCAATGGGATCTGATGGTATT
 TTCTGCTCTCAGGACTGTTTACAGTGGAACCTCGGTGAGTCCGTTTCTTT
 TTGTTTTTTTTTTCTAATTATTATTATTAGTAGTATTATAAATCAATAT
 TACTGTTGCTTATACATATTGTTGTACATTATATACATAATACATACATT
 ATATACAGTATATAGTATACAGTAGTATATAAATATTATGTATTATATATA
 TATAATGTATTATAAATATGTCATATCTAATATATGTCTGTATTAGATAT
 AATGCATATATATTATTGTACTACAGTCATATTATAAATACATTTACTTAT
 ATCTGCCTTTTTCCACACGTTTCATTGACCTGATTAAAACTAAATCCTA
 AAGGCAGAAGAAGATGAAAACCCCAAAATTAACACCAAAATAATTGCAGCT
 ATAGATCATATCTATCAAAGCAAATTTGCCTTCAGTCCACATCACGAAAT
 TAACAATAGAAAGGTTTAAATTTGGAACGTACAAACAATGACAAATAACC
 CCCAATGGCTTTTCTCTTCTTGCAAGGAGAGCGTCACACCAAAATAGGTAC
 GTGAGGTGTTTGCTACCTTCGTTTGAAGGAAGAAATTCATTAAATAAAA
 CCTCTGTCCAATATGAAGCCGGGGTCAAATTAATCATAAATCACCCTGA
 TTGTCCATGAATTAACAGGGAAAAAAGGCTAAACTTGAAAATAACATTT
 TTTTCATCTCTCTTTAAGGGGAACCTCACTGCAGAAGTTGGTAAGTCTCT
 TTCCCATCAGTTTAAGCAAAAATGGTTCATCAGATATATAAATAATCCCTT
 ATTTCTGCTTGTTTTTAGGGGACTACAACAGGAACTTCGTAAGTGCCTT
 TAACTTCTCCCATTAAGAGTTAAACCTTTCAATATTTTGTATGCTTCAAT
 GTGCTGAAGCCACCAAAAATGTGTTTAAATGTAAAGGGGCTGAGCGTCA
 AACCTGAACACTGCCATGTTGGGGGCTGAGATTCGTGGGATTTGGGTTTT
 CAGTGTGAAAATGCCTCTGGGTTTCTGTGCCTGAGCTCAGGGAAACACGA
 CCAGGGCTTCCAGTAGGAATGAGACCCCAAAATATTCTACCTGGGGCC
 TTTTCCCATTGGGAATTTATTCTGTAAATCCATATTTCTCCACGTTTGAG
 CGTCACTCATCAAATGTCACAATCTTGGCAATGTTGAGAAGATATATAGA
 TATCTATTTTAATACTGATTAATATGGAGGTGTTTGTGTTGGTCAGTGAT
 GTCATCGGGAAAAGATCTGAGTCATTGAATCCCATTTCTTTTCTCTTTA
 TTTTAAGGGAAACACGCAGCAGAACTTGGTAAGGGAAATCCCTCCCTGGG
 TTTGTTCTCTTGTCTTTCTCTTTTGGAGGGGGATTTTTCTATGTCTTCT
 TTCTATGTCTTCTTTCTATGTCTTCTTTCTATGTCTTCTTTCTATGTCTT
 CTTTCTATGTCTTCTTTCTATGTCTTCTTTCTATGTCTTCTTTCTATGTCT
 TTCTTTCTATGTCTTCTTTCTATGTCTTCTTTCTATGTCTTCTTTCTATGT
 TCTTCTTTCTATGTCTTCTTTCTATGTCTTCTTTCTATGTCTTCTTTCTA
 TGTCTTCTTTCTATGTCTTCTTTCTATGTCTTCTTTCTATGTCTTCTTTCT
 TATGTCTTCTTTCTATGTCTTCTTTCTATGTCTTCTTTCTTTCTTTCTTT
 CTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCT
 ATCACCTCAAATGAGCCTGAATGTTTGCAGTGAAGGACTGAGCACAGCTG
 GGCACATAATCATCTTTATTTCTCTCTTATTTACAGAGGAACGCGATCTG
 AAAATCAGTAAGTGCTGCCCAAGCCATAGGGCTATGCTGGGCTTCATC
 CCCACAACATGAATTTTATAAATTAATAAAATAAATAAATAAATAATTTT
 ATATTTTATGATTTTATATAGCAGTATTTAAAAAAGAATAAATAA
 CTCAGAATCTTAGGATCAATAGTAACACAATGATGCAACGTGGATACAA
 AAGCAGTAATTCCTATTTCTTTGGGTTTTATCCTTCCAGGGGAACACGA
 AGCAGAGATACGTGAGTGTTATTTTATATACTCTATAATGGAAAATTTT
 TTCTCTGTAATATAAAAATAGGCTTTATTATTTGAGGGGTTTTTTGGCTT
 AACGCAAAATCGGAAGTGCTTGAAATTTCTACGTATGAAATAGAGGATTTCC
 CATAGAGAAAAACAGCAATTTGGGGCTGGAATAAAAGTTTCATTTCTTTG
 CTGAAAAGTGAATGAAAAGGGGGGGGAAAAGAACATAAAAATGAGTTTTT
 TCCCTCATTAATCTGTATGAAATGGGTTGGGTTCTGAAATGGTGATGTC
 AACACCTCGTTTTGGGTTTCCAGCCCAACATAATATGTGTCTGTCTTTATT

FIGURE 10

SUIITE 40

FEUILLE DE REMPLACEMENT (REGLE 26)

TCCCCAGTACCCAATTATACAATGGGATTAATTACAGCCTGCCAGGAAA
 GGAGCACTGAATTTTTTCTGCGTCCATCCAGCATGAAGTCCATCAGACT
 TAAGCTTACAGCTTAAAGAATGGTTCAATTTTTTCAITTAACCCCTCGT
 AAGTTAAAAGATGGACTTCAGCATCACAGAAGTAGCCCAGAAATAGTCAA
 AAAATGGGTCAATTTCCAGAGCACCCCCCACACTTTCTTTGGTGAA
 TAGGAAAACAAATATTAATACTAATTAATTGGTTTTTTTTCTTTTAGG
 AAGATGTTTTGAGGAACACAGTAAGTGCCCTTTCTCCCTTCTTTAAGCA
 TCACTTTTCACTTTAAGTCTGCATCACAGTTAATAATCCATCTCCTTATT
 ATGCATTTTLAGGGAGAGGCGAAGAAAAGTTGGGTAAGTCATTTGGTTAA
 TTGGGTTTCTGCTTGCAGACCCCATCCAGGAGCTCATGTCTCCTCTTAG
 TGTCTGCACCTGTAGAAATATCCAGGTTAGACGTGTAGGTAGGAAATAGT
 GACCTGCGTGGAGGTATTGCAGACCCCATTTATGTGTAGGGGAAGCAGAA
 CATCAAACTATTGAGCCTTGAAGCTCCACGAAGACAAGCCACCCTCTTAGA
 TTTCAAGCGAAGTCGAGCTGAATAGATTTAATTCTTTCTTTCCCATAGTA
 AATGTGACTCTGGACCCAGAGACGGCCACCCTCGCCTCGTCTCTCCAA
 GGACCAGAAGAGCGTCCGATGGGAATACAGCCTGCAGGAATCCCCGACG
 GCCCCGAGCGCTTCGACGCCGATCCCTGCGTGTCTGGGTTGTGAAACCTTC
 ACCTCTGGGAGGCACTGCTGGGTGGTGGATCTCACAGAAGGGCAGTACTG
 CGCCGTTGGGGTCAGCAGGGAGTCCCTGCCAGGAAAGGAGCCGTCAGCT
 TTAACCCTGATGAAGGCATCTGGGCTGTGCAGCAATGGGGGTTCAAGAAC
 AGAGCCCTCACCTCCCCTCCGACCCCACTGAACCTTCCACGGGTTCCCAA
 AAAGATCCGCATCTCTCTGGACTACGAATGGGGCGAGGTGGCGTTTTTTG
 ATGTGGAGAACCAATGCCCATCTTCACTTTTCTCTGACCTCCTTTGGT
 GGGGAGCGGCTCCGGCCGTGGTTCTGGGTGGAGCTGGGCTCCCTCTCACT
 GCCCAGATAACCCCGGAATCCCTGGAGGTGCTGTGGAGGTGCCTTACAGC
 AGCTCTTCCAGACCGGGGTGGAAAAACTCTCAGGAAAAGCAGCATTAATA
 CCTCATTTCTCCCTCTTCCAGTCAACCATTGTCTGCAAAAGAAAAGGAAA
 CCCATCTCAATGTCATCAGCATCCTCCGTGTGTCTGTCTGGTGGCCCC
 CATTGATGTATGGGGTGGCTCCTGTTGGTGTCTGGTGCCCCCTATTGACG
 TATGAGGTGGCCCCCATTGACGTGAGGTGGCCCCCATTGACGTGAGGTGG
 CCCCTATTGACATATGGGGTGGCTCCTGTTGATGTCTGGTGCCCCCATT
 GACATGAGGTGGTCCCATGACCAGCCCCCTACCCTGGATCCAATGCCTCC
 TGATTGCAGTTCCAACTCTAGGGACGTTAAACGACCCACAGAGAGGATG
 GGGTCTCTTTGGTCTGATGGAGAGAGGTGGCCACCAGGGTAAGTCGCTG
 CCTACATCACCCTGGTGTCTTGTCTCAGCAGCTGGTGTAATTTCTGCC
 ATCTGGGCTATTTCTGTAGAAAGCAAAGAAGCTCTGCTGGTGGGCAGCTC
 ATCTCCAGTGTGAAAAGCAAATGCAACGCATGCACCCTGCTATCCAT
 GTGGCCATCCCTCTCCATCAGCTGTTGAAGGAGAAATCTGCACTCAGAAG
 AGATTGAATTGGGCTCAGATCTGGCTTGGGAAGATGATGATTCCAACCAG
 AGTCCAGGAGACTTTGGGGAATGCATGAATCCTATAGGAAAATGGATAAC
 CCTTCATCCAAGAGCAAGCTGGCATGATGCTCTGGGGTGAAAACCCATAA
 TGCCACCTGGTTTTAAGGTTTGGGGTGGCTTACAATGTGCAGCTCTGCTT
 CCGGCGAGGCACTGGGAGCCCTAAACCCATGGAGAGGTCAAACCAGTGCT
 GGAGGTCAATTGTGGGCCAGCTGCAATGGGAGGTAGGCAATTATGGACAT
 CGCTGAAGCCACCCACGCTCTGGGGAACCTGGGGTTTTACCTTTCACTG
 CACTTTAATGGGATTTCTCATCAATGTCTGCATGTTCTTGGCCACCTGTT
 TAAAAATATAATAATAATAATTAATCTTTTGGCCCACTGCGGGATGAGC
 AGCTGGTGGTTCCCAGCTCACAATAAACACACTTGAGACTCCCTGGAGA
 ATTCGCTTTCTTTTGCAGCTGGTTCCATGTGGGGCTGTTACAGCCCTCT
 GCAGCTCATAGGCTTTTCTTACAGCCTCTGCTCCACCTATTGCTGAAAA
 GGGGSAATTTGAGATGGATCCCATTTTGTGAACATCTCCACCTGTGGG
 TAATGCTCAGACCTCTCAGCCCTGTGGGTTTAAATTTCTTTCTGACAGT
 TAATGGGTTGGGGATGTTTCACTGCAATAATTAGTGATGGGATAGGGG
 AGGCAGGAGAGGATCCAAGCAGGGGAAAGGGAGGGGAAGGACATACTGT
 GTGTCTCTG

FIGURE 10

SUIITE 42

2044160.04400

AGAATACCATTCCTCCCTATGCCAAAGCAGAACTGAGGGGCAAAAATAGTG
 GTTATTTAAAAATATATATGTTTTAATTGACTATCAACAGGGCGAATGGC
 ACAAAGGTTGCATCACGCTGTGTGGTGGGTTTTGATGCAACTCAAAATTG
 CAACTAGGAGTTCCTGTGCTAAGTGCTAAGGAAAAATGAGATGAAAAATG
 AGATGAGAAGCCGCCCAGCTGTTTAATTAAAGCAGTTTGGTGACTGTCTG
 GCTATGGTGACTGAAGTAAGCAAGCACTGTGCTGCAAAATGCCCCATCTC
 TCCTGGAAGTCGAGGATATTTCCCTGTGGCCAATAATGACAGAGCATT
 TTAAGCCCAAATCTTTATCCAGCCAAATTCACAATGGAGACACCGATCT
 GCAGGGAGATTCCAGATAGCACATTCTGTGTTTCTTAAATAGGGCTGAT
 ATTTCCCTCTGTCCCACATGTGATCCATCTGAGCTCACACAGCCCTTAAA
 AATCTGCAAGGATCTTGTCTGCTGCCTGGCGGTGAGTGTCTTTTGGGG
 CTATTTGGCACCATTTTGGCAACGGGAGTTGGCAACGTCAACCTGAGTTC
 TTCCAGAGTTCATGAAAAGCTTTAAAATCGGGTTTTTGCACTTTTTCCA
 CGATCCACCTCTAGGGACCAAGCTGGTATTGGGGGGGGGGGGGGTGTGC
 TTTCAAGGGGGGTTTGGGTTCTGTTTTGGGGGATATTTGGCATATAAAGGT
 GGTTTTTCACCTTTGGGATGGGCCTTGGAAGGAGTGTGTTGTTGGCTGTC
 CCGTTGCCTTTCTTCTACAAAGCCTTTCTTCTACAGAGGCTGCACCAA
 GGTCTCTTTTAGCAGATCACAAAGAGAGGGGCGAAAGAACGAGGTTAGAA
 TTCAAGTTTTTAGGGTTGAAATATGGGTAGGATGATTGAGTCCTTCTCT
 TTGTCGGTACCAAGCCTGGAGATACCAATCTGAGATGTCAAACTGCACAA
 TGGAGCCTTCAGTGGGATGAACCTCAGGCCAGATGCCCAAGAGAGGTGTT
 AAGTTTTGGTTAAAGCCCACATCAGTAGCAATAGAAAAGAAATGGGTGATT
 CATCCCTTCCAAAACCTCAACTTTTGAAGAAATAGGGGAAGAAATCCCGC
 TTCCGTGCTGAACCTTCTGTATTTTCAACCCAGGGCAAGAAAGGAGGAT
 CGCGAGAGATTGGAGACCACAGCACCATGGCCTCTGCTGCTTCCAGAGCA
 AAGGAAAAGGGAGAGGGGGGCTCCCACCACCCTATCCCAGAGCATCAGAT
 GGGCAATGGATGCAGCAGCTCCGTGGGTGCTGGAGGTGGCACGTGGCAGG
 AGCGAGGACGGCTCGGAGATACCGAGGTCTCAGCCACCGAAACCATCTC
 AGGAAAGGGAATTTCCACACAAAACCTCCATTTGGAGCACCTGGCAGAGAA
 GCTGAAGCTTTTGGGGCTGGATGGAGACAGAGGGGAGAAGGAGAACTCT
 GCTCGTGGCGCAAGAGGACATTCCCCTCCAATGGACCACGGGATGATGGA
 GGTCCCCTGGAGCCCCCATAAAGGAGTCACTGCAGGAGGATGTGGTCAG
 CCTGTGTTATTCCTAAAGCCCTGTTAATCCTTCATGTCCATGCTGAA
 AACTTCTTCTGCGAAGTCCAACACATTGCATCTCTTCCCTTCTTTCTC
 CCATCAATATCTCCCCAACCCCTTTTTCTTCTCCTCCAGGAGCAGATT
 CACAGCGATCTGGAGAACCTCAAGAAACAAAAGGAGGAGCTCTTAGAACT
 CAAAAGGAGTGGGGAGAGGCGATGCCAAGACCTTCTGGTAAGAAGCTGTT
 GCCTTCAAGCTGGAAAAACAGAGGTCTTTTTGGGGTCCACGTTGTTGATT
 TTCCACAACCTACAGACACGGACGGAGGCTGAGAGGCAGAAAATTGTGTC
 AGAATTCGTCAGCTCCGCCGTTTTCTGAAGGAGAAGGAGATGGTGCTCG
 TGGCACGGCTGGGGGAGCTGGACAGGGCTGTGCTGAGGAGGCAGGAGGAG
 GAGGAGGCCAAGGTGGAGGGGGACATTTCTCTCCTCGGCATCCTCATCTG
 TGAGATGGAGGAGAACTCAAGCAACCCACACGTGGATTCCCTACAGGTTG
 GATTCCTACGGGTTGGATTCCCTATGGGTTGGATACTCCATTGGACCTCT
 CCTTCTTGTCCACCTTCTCCAAAGCTGGGGGAGATTGAACCATTTTTTC
 CTATCTCTTTCAATTCCAGGATGCCAGAAGCACGCTGAGCAGGTATGTGC
 TCCTTCAACCTCATTCAACGGGGTGGAAAGGGTCCCCATCCCCACACCC
 ACGGATTCTAGCAGAGAAATGAGAAATGCATGTGATTGAGGCAAGGTTGG
 AAGTCCCAGTGGGGGTCAAAAAGTGCCTCAGTGTAGGAATGGCCCCAAGAGA
 AAGACCTCGTGGCCATTGGGGCGACCCAAGGGACCGCATCTGTACGGAG
 CAGGTTGGCATCCCCAACCTGTGACAAAGGGACATTCTGGAGCCAACCA
 CCTCAACCTCCATCCCCACATCACCAGAGATCCCCACACTCATCACCACC
 AGTCCCAGCACTGATGAGATTGTGTCCAGGTGGGAGAGGGGCGAGAACAC
 GAGGATGACAGAGTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT
 TCCCAGCAACATAAATCCTCAGGGAGACGTTGGGGAGATTCCAAGGTAT

FIGURE 10

SUITE 43

10043160 - 01402

TGTGGACCACAGTATTCTTACCACATAGGATTTGCTTTGTACTGAAGGTT
GGGGGGTTTTTTGGTTCGTTTGAATAGGAGTTGTATACACTATTGGAAAAC
AATTTGCATTAACTCACACTATCAATCATTCTTAGGCCTAAGAGCATCTG
TTTTTTAGGACCAAATCCCACAGATCCCACATAAAATCCTGCACAGATAT
CCATGATAAACATGGTGGGAACTGAAGCAGGCAGATGTGGGACATGACAT
CCAACCTTCTGTTCATCCCCAGATCTTTTTCTATCTGAGCTGGAGAAGGA
GGAGGGAGCATCTGTAGGAGAAGAGGGAAAAGGTGAGTCCTTAAAGCATT
TTCCTTTTGCTCCATTGGTCATTTTTTTAGCCAAAATACTGCGTCAGAGC
ATCTGGAAAATGATGGTTTTTGAGCTCATTTCTGGTTTCCTAAAGGTGATA
TAAAGAAGCTTTCCTATATTTTACGCAAAGGTTTTCTGAGCTGGAAAATAT
GGAGACATCGCTGATCCCAAAGTAGATTTGGGGTGCTGTTCCAGCTTTAG
GGTGATGCTCACCCTTTCTTCTCCATCCCCAACAGCGTTTGTCACCCTG
GACCCCACTGCCCAGGGCTCGTCCTGTCCCGGGACCGACGTGG
GGTGAGATGGATGGATATGGGGCACAACATGTCCCTTGTCCCCAACGCT
TCGATGTCTCCTGCTGTGTGCTGGGCTGTGAGGCTTCACCTCAGGGTGG
CACTTTTGGGATGTGGAGGTGATGGGTGGTGCCACGTGGGCACTCGGGGT
GGCACGCAGCTCTGTGCCAGGAAGGGTTGGCTCACTTTCCACCCCGATT
ATGGGATTTGGGCTATGGGATGCTGTAGGAACAGCTTCCGAGCTTTCACA
TCTCCCCCATCC

FIGURE 10

SUITE 44

204710 0973400

82/110
D12FOR.txt

GGGGGGCGATATGGGTGGTGGGACATGAGGGGGCCGGGGGGGGTCCGGTC
TCACCCGCCAGCAGCAGCCGCAGCCCCGCAGCCATTGCTCTCCGCTGCTT
TCGCTTTCGGCTCCGCTGTGGCCCCACCCCCTCCGTCACTTCGTCAATAT
TAATTTTAAATCCCTGAAACCCATTAAAAAAGGGTCGGAGAGGGAAAAC
TCATTTCAGGAACAGTGTTGGAAGAGGGGACATGGGTGGGACAACCCGGCT
TTCCCCACAGGCCGACCTAAACACAGCCACTGCCACCCACCCCGGATCCA
TGGGTGACGTAAGGATGAGGTTCCAGCACATATTGGACCCTTCTGCGTTT
GCATGG

FIGURE 10

SUITE 45

004350 01400

2043150.01402

GTGGGGGGCAGCGTCCGCGCTGACCTCGTCTCGCTGTGTTTCAGGGCGCC
CCGTGCGCGCGCTCCAGGTAACGTCCCGTTCCCATTCCCGTTCCCGTTCC
CCGTTCCCGTTCCGCGCTGCGCGGAGCGGCCCCGATCCCGGCGCGGGGCT
CAGCTCTGCCCCGTCTCCCCGCGAGGGATGCTGAATTAGCTGCTGCCCCGCC
GAGCCGCTGCACCCGACCCCCCGCTCTCCCGGCCGTGCGCTCGGCTCTC
CCTCGGGGTGCCACCGCGTCCGTTGGAGATGTCGCCACGATGCACGCTTC
GTCCCCATCCTAATAAACGCGCTGACTTTGACCCCGCTGTTGCTGCCCG
TGAATCATTGGGGACTTTCGCTGCGCTGGGAGGAGGGGAGGGAAGTAAA
GCTTCGTGGAGAAGTAAACCCAGCACCCCTATGGGTCCCACGGGACGTGGA
TTGGTGGGGATGGGGTGGGATTGGACTCTTGGTGGTCATTTCCACCCATA
GGGAGCTGCGCGGCCACCCAGTGGTCCTCATATAGACTCCATGGTCACACC
ACTGTCACCTTTTGGTCAACCCATGATCCCTGTGTTACCCCTCCGGGGTCC
CTCAGTGGTTACCCACGTTCCCCCAGAGGCTCCTCCTGTGCGCTTCATC
ATCTCACCCCATTGACCACATACCCCCCTCCCCCTATGGATAACCCAAAG
CCATCACCACTGGTGTGGGATGCAAACACGGGGCCCCGGACCTGTCCCT
ACAAGCACAGGGTGGTGACACAGCCCAGACAGTGATGCTGTGTCATTTGT
CACCAGGCAGAGGACACACAGCCACAGCCTGGCTCAACTCGAATAATATT
TTCTTTTATTACATGTTAAAGAATCGAAAGGTTGGAAACATACAGTAAGA
TGAAAACACGGCTCTAAGGGTCTAACAGTGGGGCAGGAGGGTGGGGGGGA
GGAAAAAAGAAAAAGGGAAAGAAAAAACCAAAACAAGTAGAAAAAA
ATGATACAGTCAACGTAAAAAGGGGTGGCCCTCCCTCCCCAGTGGGAA
CATGCGGCGCTGCGTGCCGGGGGGTTTTATGTACAGGGGGCGGGCAGCTC
CAATAAATTAAACCTCCAAATACAATGAGGGGGGAAGGGGGGGTGCAGA
GCCCCCTCGCTGGGTGGTTTTCTTCTTTAAATGCTTTTTTTTTTTTTGT
AATTTTTTTTTTAATTTTTTTTTTTTTTAATTTTCTTAAAAACCCCAAACCTT
TTTCTCCCCCCCCCTTTTTTTTTTTTGGAAAAATCCCACGAGTCAG
GAGGAAAAAAGAAAAAGCCAAACCTAACACAACAACAGTAAACCT
GCTGGGGGGCACCGCCGACCCCCCTTGTCCGACCCACAGCCCCACACT
GCCCTGGGGACGCTCGGGGGCTCCGGTCACACCGGGACCCCCAGCTGAG
TCCATGGGGCGTCCCCGCGGCTGCTGGGGGGCTCTCGGTCTGCTCCATGC
CGGCCCGGTCTGTCAGAGCCGCTCGGGATGCTGCCCCATGTGGTGCTGTG
GGGTTTAAACCCGAATCCGAGTCTGCTGGTGTCCGAGGACGAGGAGCTGGAA
CTGGAGCTGCTGGAGTCCGAGCTGGAGCTGGAGGCGCTGAGCCGTGAAAC
AGCCACCTGCTGTGCTGACTCGGGCTTCTCGTTGGCTGCAATGGGACAAC
ACTGCGCTCAGCATCACACAGATCACACCCCAATCCACTCCAGACCCC
ACACTCACCTTTTTTTGGGGTTTCTTGGCTGAGTTGAGCTGCCCGTGA
CGTCTGTCAGCCGCTTCTCCAGCTCCCGCTTCTTCTCCAGCGCCAGTTCT
TCTTTCGTCTTCCCCACCGGCTTCTTCATGGCTGTGAAATTCAGGTTTCAG
CCCCACACCATCCACCTCCACCCAGGGCCGCCCCCTGAACGCAGCCCC
CCACTCACTCTCGCTATAGGGTTTGCAGGGTTTCTTCCGCAGGCAGGACA
GCACGTAGCGCTCCAGCTCACGCAGTGTGGAGGGTTTGGGGTCTCGAAG
TCGATCTCGATCTCCTCGGGGTGGAAATCACGCAGTGAGGGCTCCCGGGA
CTGGATGATGTGCACCACACGGCCAGCTTCTCCCCGGGCAGTTTGTGGA
TGTCCAGGCTCAACTGCCTCTTCTCATCGTACGTATCGGTTTGCTCTCC
TCTTCTCTTCCGAATCGTAGAGCGTGGGCGGAGGCGGCAGCGCCGCTTT
TGCTGCTTTCTTTGAGTTCTTGCAGGAAGCAAAGCACCATCAGGAAAAATG
AACCTCAGGAATCACCCACAGCTGACCATCATCCCCAAAAAACAGCCT
AGACTCACTTGGAGCTGCCCCACCGCTCCCCCGCGGCCACCTTCTTG
GCTTTGCGGAGCTGTGCCTGACGCGCCCGGCTCTCTTCATCTCCTCCTCG
CCCTTTGTGCTTCTCCGATTTCTTCTTTTTCTTTTTCTCCGCTTCTTTT
TGGGTTTGGAAACGGGGCCCTGTGAGAGGGCAGCCAGCTGCTCGTGCACG
GCCCGCAGCTGTGGGGGAGACAGGGGGTGGGGCGGGCATGGGGAGCAGG
CACAGGCAGCAGCAGCGGCCAGCTCCGGCCCTCACCTGCTCCTGCAGCT
CTGCCAGGCGGTTGGCACGTTCTCTCCGAGTCAGAGCTCTCCTCGCTG
TCTGATGAGCT

FIGURE 10

SUITE 46

10043160 011402

TTCATCTTCATCATCCTCATCGCTGGAGGATTCTCGGAGGAGGATTTGG
AGAGGGCTCCAAGCAGTGGGGCAGACACTGAGGGTGGGCTGGCGTCCTGC
GGCTCATCAGGCATCTTGGCGTAGCTGAACTCAAAGACATCCTGAGAGAG
AGGACACAGAGGGGTAAGCTGACTGGGCTGGGGGTACGGGGCTGCTGGG
TGACCCACCCACCTGCAGCTTGCGGGCCATGGCCACCACATCGTGGTCG
GGAGGGTTGTATTTGTAGCAGTTGGAGAACATTAACCGGACATCAGCGGC
AAACTCCTGTGCGTCATGGTAGTCCCGGTTCTCCATCTTCCGCTGTGGGA
AGGGAAGGCGTGAGCAGACCTCAAAGCCACCCCCACAAAGCCCCCATGA
GGCTGTGCCAAGGCCCACGGAGTCCCCAAGCGAACCTTGATGGTGGTCGAG
GTCCATGGGGTGTTTTGATGATCTCGTGGTAATCGTGCAGCCCCAGCGCCG
AGGCATCGACCGGCTTGTAAGGGCCATGCGTAGGCAGCGTGCTTCTTG
GAGAGCAGCTCCTTCAGAATCCCATTCAGTATTTGAGCTGCTCCGACAA
TTTGGCCTTTTGGAGGTCTGATGCTGCTGGGAATCCGGCAAGTCCTTCT
TGGGGGGTTTGATGGGGCGGCGCTCTCACGCCGTGCGGGAATTTGGCC
GCCTTGGCCTCCAGCAGCGTGGCTGACGGGGAGGATTACCCGCTGGTGGC
TATGATGGCGGTGGTGGTAGGGGTGGTGGTGTCTGCTTTCCGCTTCACAC
CCTTTTTCTACCAAATACAGAAAGGTTGATGAATGGGAGGCCAGCACA
GCCACAGAGCCTCCTCCCGTGAGCGAAGAGCTCCCATCTCCCACCTTGG
CCACGGGTGGGTGGGCGCAGGCGCAGTCAGCACAGCCGGGGCAGTGGAG
TGCAGCGACTTGAGGAGCGGAGCGGAGATGACGGACGGGTGGGGAATGTT
GACAATGGTGGTGGCGATGTGCGGGCTTGGGGTGTACACAGCGGTGTGGG
ACACAGAGGAGACAGCTGGCACTTGCTGAGCCGCTGTGAGACCTGCCAGG
AGCGCTGCGGACAGGCAGAACCCCCATTAGCACCAAGGTACCTTCAGTGC
TCTACCTGAAAGCGCAACCCAAAGAACCCAGGTACCTGCTGCCCGCGAC
GCTCCCTTCTTGTGGCTGTTTTTGGCCACTGGGACCACGATCTCCTGCTC
TTCTGGTGGCATTGTTGGGCCACCTTCTGCAGGAAGATCTTCTCCAGGGTTT
GGGCCATCAGCACAAATGTCATCTGTGGGCTACAGGGACAACCGAAACGTC
ACAGGATGCAGAGATGGCATCAAAGGCCTCAAAGCATCCATGCTGCAGTC
CTCACCTTGTTATAGATATAGCAGTTTGTGAACATGGTGGTGAAGTCCTG
CATGCACTCAGCTGCCCCCAGTAGTAGTTGTTCTCCAAGCGCCGTTTGA
TCGTCCCCATGTCCATGGGCTGCTTGATGATCTTGTGGTAATCCTGCATA
GGGGATGGACAGTCAGCGCCGTGTTGGTAACCACACTGCACCCCTCCAG
CCCCAGAAGCAGTGGTTTGGGGTTTTTAGGAGCTCAACATCCCCCAAAGT
ATCAGGACGTTGACACGCACACAGATCCGCTCTCGCACCATGCATCAAAA
GCAGGGCAAAAGGGTGCAAAGGGATGAAAAAACACCTCCGGGTCTGGTCC
CCGCCCCGAGAGTGCCACCGTGCTGCTCTGTAGGGGACCTTCAGGTGCTCT
TGTTGGTTGCCATACGCTATAGGGACAGCCAAAACACTGCTGTCCACAGCA
TGAGGTTGCAATGGGGGCCACTAATGCTAAAGTAAGAGCAAACCTATGTGG
AATTTACCTCTGGGCTTTAAATCCTTGGGCGCCACAGGTACACAGGGGGC
TGGCGTTTAATATTGGGGTCACAAGATGTCTTCTACAAATTTCATGGATGG
GAATCTGCAAAACGCATTTCAGGGCACAAGAGATTAGGTGAGGAAACATCC
GGGTTCCCTCTAGAGCAGCTGCGTCACCTCACCCATACCCGTGCGGTGGC
ACTGGGAGGGGACAGCAGCTCTGAGGACATCAGGTACCTACTGGGGGGG
CTTCAGAGCCTGTGGAGTTGGGATTATGCCCCCTAAGAGAGGGGCGAGGCCA
GCACAGCCCAGGCACCTGCAGCTGCATCTCTGTGGTGGAGCCCATAGAGG
GGACAATGCTGTCCCTGTGGCACTCTCAGGCTGGGGACCACGGCTCGGGG
TGGCCCTCAGCACCCAGGGGACAAGTCTGGGGACACACAGCCATGCTGGG
GGACCCACAGGAGGGGACACGTTACCGGCAGACCCAGCTTGACGGCGTCG
ACGGGCTGACGGAAAGGCCAAGCGAACTGGTGCTTCCACAGGGCTTTTCAT
CACCCTTTGTGCAGGTACTGCAGCTGGTGGTGACCCGGCCGGGCTTTT
TGGGGTTCAGACCTCTGGCGGAGGGGGGTTGCTGCTGGGGGTCTGTAGG
GCCGGCACCGAGGCCATGGTGGGGCTCTCGAAGCCCTCGTAAAGCAGCGA
GGGTTTGGCGATGCGTTTGCCCGGGGTGATTCGCTCGCCAAACCCATAA
GCCCGGCATTTCCCTCCCCCAGAATCCTGCAAGGGAGCAAAAGACAACAT
CAGCAAGGATGGGGCCAGCGTTCCACCACCAAGGTGCACAAGAACAGCTC

FIGURE 10
SUITE 47

AAAAAAAGGCCAAAAAAGTAATCAAAAAAAGGAAGGTTGGAGCAAACAAA
 GAGTCAGTGCAGGGGGTGACATCAGGGCCCAGCAGTTTCACACCTCGGG
 GTACGACAGCCTGCACTACAGCATGACAAGGCAGCACCCAAACACTGTGG
 CCTCAGCTGGATACACAACAGTGGGCTCCAAATGTCTGGGGACGGGGGC
 AGAATTATTTAAGTGGGAAAATGAGGATTTAAGCAGCTGGGAGAGGTGG
 GATGTCTGCAGCGTGAGGAGAATTTGTACCGGGAAAATACGGTGAATGT
 CGAGCACTGGGGCTGCTTTCTCAGGCAGCTCCCAGGGTGTTCCTCATCCT
 GCCAAGGACGTGGTGGGAATGACAAGGAAGGAAGGTGACAGAAGGACACA
 GCGGCCCCAGTAGTGGCGGTACAGGGTGGGAGGACACGGTGAGACCCCTC
 AGCATGCTGACAGTGTCCCCGAAAGCAGCTCAGTCAGCAGAGGTGGCAGC
 AGGGCCCTAAGGGCCCTTGTGATGCTGACCCCAAGGACCAGGGGTATGAG
 GAGTGGATAAATGGGGGTGGCCCAGACAGGATCCATGGGAAAACAGGGCT
 GCCAGGTTCCCTGTAGGATCTGTGTCCCTGCATCCCTGACAGAATTCACA
 TGGACCACGGGGCTGCCGAGTCCCAACATCCCTGAAGGACCCACAGAAAT
 GGGAAAGTGGATAAATGGGAACAAGCAGCAGATCAATGGGACTCAGTGACC
 CCAAACCTCAGAGCTCTGTGACAGAAAAGCCCCATAACTCTGGTGGACATC
 CACACTGCACCCTAATCCCTGGGCAATGAAGGGATAGCAGCAGGGAACCA
 CTGTGTCCCTGTATCTCTGACCCCAAAGAATCCATGGAGATGGGGAATGG
 ATAAATAGGGATGGCTCTGTAGAATCCGGGTCCCATTCCCCTCAAATAAT
 CCATGGGAATGGCACTGTTGGATGCATGGCCTTGAGTCCCTGTCCCTAAA
 AATCTGTAGGAATGACTCTGTGCTATGCACCTCCCCGTGTCCCTGTTAGG
 ATCCATGGGGACAGCAGGCTGCCAGGTCCCCTGTATGATCCACAGCCCTA
 AAAGCAGCTTGGTCAACAAATGGGAGGGAACAGCGGGTCCCTAAAGAGCG
 CCAGGTCCGCATGTCCCTGTCCCCAAAGGACCCACAGGTACAAGGAACGA
 ATAAACAGAGACAAGGAGCACTCAGTGGGATACAACTGATGTCAGGTGCA
 GAGCCTTTGAACACAGAAGCCCCATCTCCCCATAGGATTGAGTCCCCAT
 GCCCCTGTTGGAACCATGGGACAGGGAGGCTGATGGATTCCCTGCAGGA
 CTGAGTTCCCTGTGTCCCTGACCCCGAAGAATCGATGGGGACAGAGAGTAG
 ATAAACAGCGATAGCCCTATAAGATCCAGGTCCCCGCGTCCCTGTCAGGA
 TCCGTGGGGACCGTGGGGCTGCCAGGTCCCCACGTCCCCGTCCCCAAGCA
 ATCGATCCACGGGGATGAGGAACACATAAACGAGGACAACCGGCATACAA
 ACGAGATCCAACCGGCCCCGGGTGGAGCACCGGGACGCGGCAGCCCCATA
 CCGCCGTCCCCGCAGCAACGCCATCCCCGGTTCATAACTGCCAACACCCC
 ACAGCCCCCCCCCGGCCCCCATTCCCTGCCCTCATCACCTACTTGCTCTGG
 GGATTACATTCTGCAGCATGCCGGCGGCTGCGTGCCCGGCCCTGGCTCC
 CGGCCTTCCCTCCTCCACCTCCTCCTCCCGCCGCGCCTCCTCCGACGTCC
 CCCCCACTTTGCCACCGAGCAGCGCCCGTTAAGGCAGCGGCCCTCGGGC
 GGGCATGAGGCGGCGGCTCCGGCCGGGCCCCGCGCGCGCCCTCACATCA
 GCGGAGAAAATGGCGGCGGGGCTGGATGGAGAGGGGGACCTTCCTGCT
 CTCCGCTGCGCACAGAACCCGCGCGACGCCGCGGATATAGAGCCGGGAA
 AGCCGGTAGGAACCGGATAGATCCTCGGAAGGACGGTGTGAGGCGGATGG
 AAGGCGGACAGAGGGCGGATGGAGGCGGATGGTTTCAGCGGGAGGGCTCCA
 TCTTGGCTCGTAGGCCCCGAAGAGGAATCGGTGCCGGCGGCGCAGGCAG
 GGGTCCGTACGGAGGCCGGGGAGGGTCCGGTGGAGCCGTCCGGGAGCGCG
 AAGGCGGGGGCTGGGCCCCGCCGGTGGAGGATGGAGGCGGATTGGGGCCG
 CCCCCAGCGCGCGCGAGCCCGACCCGCACCGTCCCTTCGTCCCCACGA
 AATGGCGCGCTCGGCCTGCCCCGGCGGCCCTTATATAGACACCACCTG
 GGTGCTGATTGGTGGGTGGACGCGCTGACGTGAGCCACCCGCTTGACCC
 GCCCTGCCGCTGCCTCATTGGACGGCGGTGCTCACCGCGCAGCGCTCCTC
 TTGGCCGCCCCGCACGCCACTCACCCGCGCGCTCCCCCCCCGCCCTTCC
 GCCCGGTACTGCGACGGTCATTGGTCCGTGCTGCCATTCCCGGCGCGGCG
 ATTGGCCACCACGGATCACGTGAGGGCGGCGCTGTTGGCTGTTGTTGTGG
 CGCGCAAGGTGCAGAGGGAGGGGGAGGGCTAAGGCGGGCGTCCGCATTT
 TGTGTGGCGGCGGCCAACGGGCGCGGGCTCCCGGGGGTTCGGACCCCTCC
 GCCCCAAGGCTCCCTCAGGGGCTCAGGACAGCCCTGCTCAGTCCCTACGGG

FIGURE 10

SUIITE 48

204770 "03E402"

GCTCGTGAGGCTCTCCAGCCGACAGCGGCCATCGGGCAGCGGAGCCGCG
GCGGTCCGCGAGCCGCGGGCTGAGCTGTGGTGAGGTAGTGAGCTGGGTCC
CGGGGATCCTGAAGGGTCTGAGGTAATCGCGGCCCTCAGCGCGGTCCCG
GGGCCTTCAGTGCCACCCACGGTGGTACTGGGGCCCTCAGACCGTCCCC
TCCCCACTGCCACGGCGATCCGGAGGGGGGGGGTCCGAGGCCGCCCGT
GTCTATTTCGAGGTGCTCTGTGCTCTTCTTCCCCACGGCAATTCTGGAGG
GCTCACAGCTACTCCAGAGCAGCCCCATAACCGTCTGGGGGCTCACTA
CCACCCTACAGCAACTCAGAGCCTCCCCCACCACCCCAAAAAACAATC
CTGGAATCCCCAAGGCCATCCACACCAACGCTGAAGGTCTCAAAGCCCC
CCCTCCCCACACACACCGGTTCTAGGGTCTCAGAACCACCCACAGCA
ATCCTGGGACGTTCCACAGCCCCCTCCGTAGTAATCTTTTAGTTTCTCAAG
GCCAACCCGTAGCACGGGGGGCCTCCGCTGCCTCCCTCGTGGCAATCCT
GGGGGGCTCAGTGCCACCTCACAGGAATTTGGGGTGGCTCAGAACCCTG
CAAAGCAATCCTGGGGTCTCGAGGCCACCCACACCGATCTCAGGGTGC
TCAATGCCACCCACAGCTGCCCCCGGGGCGGTCACAATCACCCACACC
AATCCTGAGAACTCAGTGCCACCCACAGCCAATCCCGGGGTGCCCCAT
TGCTCTCTAAAGCCTCCACCCCAATCCGGGGGTGTTCAATGCCACCCCA
CAACCCCTCAAAGCACTCCTGGATACCCACGGACACCCCAACGCCCT
AAAACAAATCCCAGGGCGCTCAATGGAACCTCCCGCGCAACCTCGGGCTG
CCCCACGCCCCCTCCAATCAAGCACGACCCAGAGACCCCCCTTTTCTCC
CAACCCCTCCGGCCCCACACGCCAAAGGCTCTCAAAGACCCCCCCCCAGC
GATCCCGGACCCGAACAGGGCTTTGGGGTCCCCCCCACGGCGCTCCCGGT
GCCGCCCCCCCCCGCCCCGTGACACAGCACTTTGGATCCCCCGCGGGCCCT
CCCCGCGCGCGCCCCCGCGCGAACACCCAAACATGGCGCTTTTCGCCCCA
AAAGCGCCGGGCACAAAGCGGCGCGCCCATTTGGTCTGCTGCCCCCGCTC
CTCGCTTCCCATTTGGCCCCCTTCGACGGCGGAGGGGCGGAACCAGATTTGA
TGGACAGCTCATGCTCACGTGTCTCCCCCCCCCGATTGGGTCTTTT
GGTTAAAAAAATAAAATAAAATCATAAAAAAGGGCGAAGTTGCCCCATC
GTCACTCACCTGAGCGCTCCCACGCAGGGCCACGACCCCAACCCGATA
TCATCCTCGCGTCGCCCCCTTAAAGCCCCGTTTTTGGGGCAAAAAATCAA
AAAAAACATCCCAGGGCAGAAAAAGGAGCCACGCGCTACGTACGTGCAC
CGTGATTGGCCACCCGCGCTCACGTGACGGCCCCGCGCCACTCCGACGGC
CCCATTATGAGGCCCGGGCGGCTCCGCGGGGTCTATCGCCGCTCCGGA
GGGGGTGATGGCGCGCATGCGCAGTGACGGGGCATGTGGTGGGGGGAGG
GAGGGGTGGGGCTGTGGGGATGCCCGGTGTTGCTGGGGGGCTGCTGTAGG
GTGTCATGGCATTGCGAGGATGCAGCCATGAAGATTACGGCATTGTAAG
TGTGCATCTGTAGGGGCCCTGGCATTGCAAGTGTGCACCTATGGGAGTG
CCCGGCATTGCAAGGGTGCACCTCTGGACGCGTTTGTATTGCAAAGGGT
GCAGCTGTGGTGGTGAATGGCATTGCAAGGGTGTATCTATGGGAGTGTAC
AGCATTGCAAGGGCGCACCTATGGGTGTGCTGGCATTGCATGCATGCAC
GTGTGGGGATGTATGGCACTGGGGGGGTGCACCAGTGGGGGTGCTTGGGA
TTGCAAGGGTGAGCCTATAGCAGTGCCTGGCATTGCAGGGTGCACGCAG
GGATGCGTACGGCATTGCAGGGGTGCAGCTGCCGGCATTGCAGAGGGCCG
AACCCGCCCCGTACGGTTGTGCAGCGCTTCCAGCTCGGAGGGCGCATTGCA
GTGCGGTGCATTGCAGTGCGAGGAGCCACTGCTGCAGGGTGTACAGTGCA
CGCCCCGAGGATGTCCCCTCGGCTCCAAACCCCAAAACCCACGCTTATT
ACCCCCCAAAACATACTTTTACACACAAGACACATTTTACCATCAAAC
CTCACGCTTTTCCCCCAAATCCCTCACAAAAACAAATCCGCGCCGTGA
TGAGACACCCACAGAGATCTACGGAGCCTACTCGTCCCCTGCTTCATTAAT
TAGAGCTGCTTATTAATTGCTTGGGGTGGCTCAGCGCCTATTACAGCGT
CGGGGCTCCCCGCTAGTTTCTTCTATCTAGTAACAAGTGACGCAAGGTAA
CTGCGGAGCGCGGCCATTGGTTGAGCCGCACGATCATCTCCTGTACAGC
GCTGGTGTTCGCCGAGATCTGTTCTGCCTAGCAACCGATGACGCGTAAA
GCCGCGAGGCACGGCCATTGGCTAAACTGGTTGCCGGTAGCAGAGGGATG
GGGGCTGCGAGCGGGCGGGGGGCTGCTGGCAGCGCTGCTGGGGCGGGC

FEUILLE DE REMPLACEMENT (REGLE 26)

FIGURE 10
SUIITE 49

CCTGGGGAGCGTCAGAGCCGGTAGGGGACGAGGGCGGGGGCGGTATGGG
 TGGGCACGGGGTASTGCCAGGGGTGTCCAAGATGTGTGCATGGGGAGTGC
 AAGGGGCTGTGCAAGGAGTGAATGATGCACTGGGGCAAGGGGTGGGCATG
 CACTGGGGCAAGAAGTTGTGCAAGGGGTGTTTGTGCATTGATGCAAGGGG
 AGGATGAGCAGGACTGTGTTTGCATGCATGCAGGGGGTGTGCATGGTGT
 GATTAGTGCATTAGTGCAGCGGGGTGAGTGTGAGATGTGGAGTGTGTGC
 AGTTGTGAAAGGGTTGCCATGCACGAGCTGATGTGCGCTCAGTGAGCGTG
 CACGCAGCCTGCAGAGTGGGTACACCTGCAACGAGCATGCATGCAGCAGA
 TGTGTCCATGCACAGTGTGTGTGTGTGCCTGGTGTGTCCCATGTCCAC
 GCCCTGTCTGTGCTGCAGAGCCATCGCTGCACACTCTGTCTGAGGTGCT
 CTTCTGCCAGCCGGACACGCCGTGCTGGGGCTGTGAGTGGCCTTCGACT
 CAGAGCAGCTCTTCTCATTGATGTCCCCAACTCGCAGTGGCTGCCGCGAG
 CTCCCCGATGGCCCCCTCGTGGCCCGCAGACATCGAGCAGCCCCACGAGCT
 GCTGCACGACGCGCGCTGTGCCGTGAGCTGCTCGATTGCTCACCAGAA
 TCGCCACCGGGCCAAACCAATGCCTGAAGCCAAGGGTGGGTGCTGCTGT
 CCCCGCTATGACCCCACTGATGGGTCCCCAGCCGTGTGTTCCAGTGATG
 CTGACCCCAATGGACATCCCCAGTTGATGCATCCCCATTGATGCATCCCC
 CACAGACATCCCCATTGATGCTGTCCCCATTGATGTGTTCTCAGTGGATA
 TCCCCAATTGATCCTGTTCCCAATGATGCTGTCCCCAATGGACATCCCCA
 TTGATGCTGTCTTGTGAAGTTGTCTCGATTGATGCATTCCCATTGATG
 TGTTCCCAATGGACATCCCCAGCTGATGCTGTTCCCACTGATCCTGTCCC
 CATTGATGCATCCCCAATTGTTTATTCCCCATTGATTATTCCCCATGG
 ATGTCCCCACTGATGCTATCCCCAGTAATGCTGTCCCCACTGATGCTGTC
 CCAATGATCCTGTCCCCAGTGTGTGTTCTTAATGGACATCCCCAACT
 GATGCTATCCCCAACGATGTGTCTCACTGATGTGTCCCCAGTCCATGTG
 GTTCCCAGTGATGTGTCCCCAACATATGACCTCACTGATGTCTCCCCAG
 TTGATGCAATCCCCAATGATGCATCCCCAACATGCATTCCCAATGATAT
 TTCTCAATATGATGCTGTCCCCAATGATGCATTCCCCATTAACGCACTC
 CCACCGACGCATTCCACCGGATGTGTCCCCACTGATGCGTCCCCACTGAT
 GTGTCCCCACTGATGTCCCCCCCCACAGGCATCCCGGTGGCCGACGTCTT
 CCTGCAGCAGCCTCTGCAGCTCGGCTACCCCCAACCTCTGATCTGTATGG
 TGGGCAACATCTTCCCCCAGCCATCACTATCAGCTGGCAGCGGGATGGC
 ATCCCCGTCAACGATGGCGTCACCCACCTCACCTACACCCCCACCGAGGA
 CCTGGGCTTCATGCGCTTCTCCTACCTGGCGGTGACACCGCACTCTGGTG
 ACATCTATGCCTGCATTGTCAACCGCGAGAGGGACAACATCTCTGTGGTG
 GCTTACTGGGGTGAGTGGGGATGTGGGGGTGATGCTTTGTGTCCCCGCGAG
 CGGTGGCTGATGGGGGCGGGTGGTGGGAGCAATGCTTTGTGTCCCTGCAG
 TGCCACAGGACCCCATCCCTTCGGACGTGTTGGCCACGGCGGTGTGCGGC
 GCAGTGACGGCGCTGGGCATCCTGCTGGCACTGCTGGGTTTGGGGCTGCT
 GCTGTCCGCCCCCGCGCAGTATGTGGGGACAATGGAGACAGCAGGGAC
 ACCCGCCCCGTACTCACTGATGTCCCCCATAAGTTGATCCCTCGGTGTGG
 GAACGGTGATGGTGATGTAATTAAAGCCCTTCATTTGCAGCGCGGTGTCC
 TTGTTTGTCCCCACTCCGGGAAGGGTGGCAATTAATGGGGTTGGGCATTG
 TCCCCATGGCCCCAGGTGGCAAAGTCTGATCCCATTGCACCGCCCATGGG
 GTGACGTGAGGAGGAGGAGGAGGAGGAGGAGAAGGAGTAGGAGGAGGTGA
 AGGTGGCACGGAGGGGATGAAGGCAGCAGTGCTCCTGATGGGGCCAAGGG
 GTTTCAGGGTGCTGGGGGCGATGGGGCTGGTGCTGAGCTGCGGGACAGCA
 GGTAGGATGTGGGGAAGTGTGGGGTTTTTGGGGTGAAGCATGGGGGTTTT
 AGGGTGCAACATGGGCTTTCTGGGGTGCAACATGTGGGTTTGGGGGTGCA
 GCATGGGGGTTTTAGGGATGCAGTGTGAGTTTTTAGGGTGCAAAATGGAG
 TTTTTGGGGTGCAACGTGGGTTTTGGGGGTGCAGTATAAGTTTTTAGGGTG
 CAACATGGGGTTTTATGGTACAGCGGGGGCTTTGGAGTGCAGCATGGGGT
 GCTGCATGTATGCATAGTGCACAACATGGGGTTCTTGGTGTGCAGTGTGA
 GTTTTTAGGGTACAGAGT
 GTGCAGAATGAATTATTAGGGTACAACATGGGGTTTTAGGGTGCGGCACA

FIGURE 10

SUITE 50

204750-01400

FEUILLE DE REMPLACEMENT REGLE 26

SUITE 51

GATGTTATGTCCCCATGGGGACCTCTGGGGGCTCCAAATGGGGATGAGGT
 CGCTGCCAGCACTGCCATCTCCCCTCTGTCCCCCAATGCAGGTGCCTTC
 ATGGTGATGTGGCCAACTCCTGCTCACTGGCAGCCAATGGGCTCTCTGCG
 GGGCTTCGACCTCACCGTGGCCTTCAACAAGAACCCTCTGGTGTGCTACG
 ACCCCGATGGCCACCTCTTCAACGCCTGCGACTGGGGGCTGCTGCACGGC
 GTGGCTGGACAGATTGCCATTGCCCTCAACAATGACAGCACCTGGGTGCA
 GCGTGCAGAGGCACGGAGACGGGCGTGCAGCAAACTGGCTGCACAGTTCT
 GGGCACAGACGGCGCTGCGCAGGAGTGAGCATTGCAAATGGGGCTGTTGC
 ACGGGGCGTTGCGTGGGGATGATGTTGCATGGGGCATTGCATGGAGATGA
 TGTTCATGGGGTGTTCATGGGGACATTGCATGGGGCATTGCATGAAGA
 TGGTGTGCATAGGGCGTGCATGGGGATGTTGCATGAAGATGTGTAGCA
 ATGATGCATGGGGCATTCCATGGGGACGCTGCATGAGGGTGTGTTTAGC
 AATGATGCATGGGGTGTGCATGGGGATGTTGCATGGAGATATTGCATGG
 GGCATTGCATGGGGTGTGATCCATGCAGCGTTACATGGGGTTCTCAAGCAG
 GGGGATGTTGCATGGAAAAGTTGCATGGAAAATTGCACAGAGGTGTGCA
 AAGCATATGCATGGGGATGTTGTATGGAGGATTGGACGGTGGCTTTGAAG
 AACATTCTGCATGGGGCATTGCTTAAGGGTCCCAAGCATGGGGATGCTGC
 AAGGAAATGCTGCTGCTTGGTGGCCTTGCAGAGTGTGTTGCATGGAGTTT
 GCTTCAAGGAGATGTTGCATGGCATATCATCTGCAGTTTTGCAGAGCACA
 TTGCTATGCACATTGCACACTGCACAGAGCAGTGCCTGGGCATCTCCCA
 GCGTGTGGCACAACGCTGTTGCAAAGGACATCCACGAGGTGTTGCAGCA
 AACATGCGCAGAGCTTGCACAGAACGTGGGATATCCCATGGGGATGTGG
 CACAGAGCATTGCGTGGGGAATCCTACAGGGAAGTGAGATGGGGAAGTTG
 CACAGAGCGTTGCAAGGGGTATTGCACAGAGGGAACCTGCAGAGAATGGG
 GCAGGAACCGTCCCCATCCCCTGCTGCTCACCATCCCTGTCCCCACTCCA
 GCTCAGCCCCAGGTCCGCATCGTCCCCGCACAGACAGGGAACCCAGCGT
 GCCCATCCGCCTCACCTGCCACGTGTGGGGCTTCTACCCCCCGAGGTGA
 CCATCATCTGGCTGCACAATGGGGACATCGTGGGACCTGGAGACCCTCA
 CCCATGTTTGCCATCCCCAATGGGAACTGGACCTACCAGACACAGGTGGC
 CCTCTCGGTGGCCCCAGAGGTGGGGGACACCTACACGTGCTCGGTGCAGC
 ATGCTAGCTTGAGGAGCCCCCTCCTGGAGGACTGGCGTGAGTTGGGATCA
 AGGGGGTGACACAGGGACAGCGGTGTCCCTGCTGTGCTCACTGCTGGCTGT
 GTCCCTGCAGGTCTGGGGCTGACGCTGGAGGTGACGCTGATGGTGGCTGT
 GGCCACTGTAGTGATGGTGTGGGGCTCAGCTTGCTCTTCATTGGTGTCT
 ACTGCTGGCGGGCCCAACCCCTGCCCCAGGTGGGTGCTTGAGAGGGACC
 CTATGGGGCTCCATGGACCTCTAAGGGGTCTCTGTCTGGTTCTATGGGT
 CTCTGGGTGCTGTGAATCTTTCTTTCTCTGTGGGTCCGTCTGGGGTAT
 CTGTTGATCCCTATGGGTGCTGTGGGGCTCTGTGGGTCTCTATGGGTC
 CTTCTGTTGGCCTCTGTGAGGTCTCTATTTGTCTCTATGCATCCCTTTGG
 ATCTCTATGGGGTCTCTGCGGGTCATTACGTGTCTCTATGGGATGTGACC
 ATTTTTGACAAGAACCCCACTCACCCCTCCTATTCCCCAACAGGTTACG
 CCCCGCTTCCCGGTCACAACTACCCCTCAGGTAACAGTGTCCCCAACTG
 TCCCTGTCCCATTTGCCATCAATGAGGGCTGAGTGACCCCATCTCTCACC
 CCATGTCCCTGCAGGCAGCATCTGATGGACACCTTCTGTACCAACTGTCTC
 COTGCGTGTCCCATCCCTGACTCTGCGCCGTGGTGTGACATTAAAGAC
 ACTCTGCAGCCTCTGTTGGTGTCTCTGTGGGCTTTTGGGGTGGGGTGGTG
 TCACCGGGGAGAGGTTGGGTTGGGGTCATTGCATCCATGATGGTGATGGT
 GATTGACATTGTGCACAGGGAGATGTCCAGGCGCCTGTGGGGTCTGTGTT
 TTAGGGCCAGTTCTGCTCAGTGCCTCCGTAAGTGATCTGGATAGGTCGTC
 AGTCATCCTAATTAAGGAGGGGACAACAGTGAATGGGGAGGAGCCGATGA
 CTCAGGCTGGGAGTGGTGATCCCAGAGGTTTCTCTGCTGTGAGTGACTC
 CGTGCTTTGCTTTGCTTTCAACAACCTGAGGGAGCGCATTCTGCCTGGCG
 CCCGATGACGTACATAAACCCCCGACTGCCATTGGCGGAGAGGCGACGG
 AGGAGCCAATGGGGGCGCGGGGCGGGGCGGAGGAGTACGAAAAGCTGAAG
 GAGCTGCCCTGGTGGGGGCGGACTTCAAGTGCAGCGGTGTGAGGCGATG

FIGURE 10

SUITE 52

GGGCCGTGCGGGGCGCTGGGCCTGGGGCTGCTGCTCGCCGCCGTGTGCGG
 GCGGGCGGCGGCTGAGTGCGGCCGACCGGGACCCCTCCCCGCCCGTAAC
 CCCACCCCGGGGCTGTGCCCGTGGGATCCTCAGACCCCCACCCGCGGCTC
 ACGGCCTCGCTGCGCTCCGCCCCCGCAGAGCTCCATTCCCTGCGGTACGT
 CCATACGGCGATGACGGATCCCGGCCCGGGCTGCCGTGGTTTCGTGGACG
 TGGGGTACGTGGACGGGGAACCTCTTCGTGCACTACAACAGCACCGCGCGG
 AGGTACGTGCCCCGCACCGAGTGGATGGCGGCCAACACGGACCAGCAGTA
 CTGGGATGGACAGACGAGATCGGACAGGGCAATGAGCGGAGTGTGGAAG
 TGAGCTTGAACACACTGCAGGAACGATACAACCAGACCGGCGGTGAGCAC
 GGCGGGGGCGCGGCTCCGTGGGTGTGGGATGGGCTCCATGGCGCAGTGC
 CGCCACACCCCCAGGCCTGGCCCTGCCCGGCGGCACCGTCCCGGGGCT
 GCCCGTCACAGCCCCACCGCGCTCGGGGTGCCGCTCCCGGGGGGACCCC
 AACCCATCCCCGCTGCAGTGGGAGCCCCGGAGCCGGAGGGGCCCCCTCACC
 CCCTGCCCGGCTGTGTTTCAGGGTCTCACACGGTGCAGCTGATGTACGGC
 TGTGACATCCTCGAGGATGGCACCATCCGGGGGTATCATCAGACAGCCTA
 CGATGGGAGAGACTTCATTGCCTTCGACAAAGGCACGATGACGTTCACTG
 CGGCAGTTCAGAGGCAGTTCACCAAGAGGAAATGGGAGGAAGGAGGT
 GTTGCTGAGAGGTGGAAGAGTTACCTGGAGGAAACCTGCGTGGAGGGGCT
 GCGGAGATATGTGGAATACGGGAAGGCTGAGCTGGGCAGGAGAGGTGAGC
 GGGGTGCGGGTGGGGGGGGGGGGGGCGGACGCAGTGTGGGGCTGGACGT
 GGGGCGGGGGCTCATCGTGGGAGCTCAGCCCGGCCCTCACTGCCGCCCA
 CCCACAGAGCGGCCTGAGGTGCGAGTGTGGGGGAAGGAGGCTGACGGGAT
 CTGACCTTGTCTGCCGCGCTCACGGCTTCTACCCGCGGCCCATCGCCG
 TCAGCTGGCTGAAGGACGGCGCGGTGCGGGGCCAGGACGCCCAGTCGGGG
 GGCATCGTGCCCAACGGCGACGGCACCTACCACACCTGGGTCAACATCGA
 TGCGCAGCCGGGGGACGGGGACAAGTACCAGTGCCGCGTGGAGCACGCCA
 GCCTGCCCCAGCCCGGCCTCTACTCGTGGGGTGAAGTGGGGGATGTGGGG
 CTGGGGGGCTGCGGGCTGCCCTTCCCCCTGCTGATGGCCCCGCTCTCCCC
 CAGAGCCGCCACAGCCCAACCTGGTGCCCATCGTGGCGGGGGTGGCCGTC
 GCCATTGTGGCCATCGCCATCGTGGTTGGTGTGGATTTCATCATCTACAG
 ACGCCACGCAGGTAAAAGCAGAGGGGTGCAGGCGGGCAGTGGGGGCTGTA
 GGGGGATCTGGGTCCCCCTTGGGAGCCCCAACCTGGCTGTGATGTGAAC
 CTGTGATGAAGCATCTCTGTCTGCAGGGAAGAAGGGGAAGGGCTACAA
 CATCGCGCCCGGTGAGTGAAGTGGGGCAGCGCTGTCCCCACCTCTGCCCA
 GTGCCAGGGTGGTCCCTGGGGTCCCTGCTTTCTCCCAAGGTACCCATTCT
 GGTGCTTGGGGCTGCTCCATGCCCATAGGGAGCACAGGGCTGGATCTCA
 CAGCTGTTCTCTTATAGACAGGGAAGGTGGATCCAGCAGCTCGAGCA
 CAGGTGCGGTGTGGGGCTGTGGGTGGGAGGGGTCCGTGTGCTCTCTGTG
 GTACTGCCAGGGCTGGGCTATGCTGGGGCTCTGCGGGGAGACCCCCGGA
 GCAGAGGGTTGGGATGTGAACCTGGCCCCGTGGGACATCATCCCTTCTCA
 TCCCCACAGGGAGCAACCCCGCCATCTGAGTGCTGTGCTTCAGCCTGCAA
 GGAGCCAACAGTCCACACCAGCATTGTTGGGGTGGTGTGAGACACAGCCCC
 ATCCTCCTGACCTCTCACATCTCATTCTGCTTCTATGCTGACTGTTATG
 CTTTGCCCTGCACTGCTTCTGTGAAATAAAATGATGGGCCATTCTGTGCT
 CAGCTTGCCCTGCATTCTGCACTGTGCTGTGGTTGGGGATGGGGTGGGTGA
 GAGGACCGTGTCCAGTTTGGCTGCTCAGGGTGCAGATGTGGCCCTGTGC
 TGAGTACCCACAGCCCTCCCCCCTATCTGCTGCTGCTCACTCCCCCTT
 CTGTACCCCATCCCTTCTCACCTCTCCTCTGTGACCCCATGCTGGTGGT
 TGCTTGCTCCCTGTCTGGCAGAACTCTCATTTCCTCAATGGCATCCCTG
 GGTGTTGGGATGTGGTCTCCTTGGTCTCCCCCAGCAGTCACTGCACAT
 ATCCACCCCACTTCCCCCCCCAGGTTGTTGTCCACAGCACTCCTATTTC
 CTCTCCCCCCCCCCCCCCCCCCCCCGCCCATCCAGCTGCCTCTGCAATC
 CTCACCTTGGCCAGACACAACCTTGGCGCACTCCACCTCCCTCATCCCGC
 CCTTCCCCCAGCTCTCCTGTCCCTGCTGGCCCCCTCCCCCCCCCCCCATT
 GTACCTT

 FIGURE 10
 SUITE 53

10043460 011402

FEUILLE DE REMPLACEMENT (REGLE 25)

CCTGGTTTATTCCCCCGATTGTGTTGTTGGGGGCTCCGCTCTTCAC
CCTGGGGGGAAGGGGCTCTGGGGGCTCCCTCATTCTCCCTGCACTTCTTAC
AGCACCGGGACTCCCGCGCTGAGATCCCATCACACCCGGGTACAAACATG
CGGCTTTATTCCAGTTCTGTGTCCACCCCCGGCCCTGGTGGCACTCAG
TGGCACCGCAGTCCATGCAGTGGCCGTTGTGTGTCGTACAGCAGCGGTAC
CGCAGCGCGCCCGGCTCGGCATCCATGTGCCACGGCACAGCTCTTGTGG
TCCCTTGTGCTGCCCCGGTGTCCCCACCTCCACCCTCAGTGTCCCCAAC
GCGCAGTGTCCCCATCCACATCTCACCACCCCTGCCCTCCCCATCCAT
AACTCCCGCTATCCCTCTGTCCCCCTCCCCCGGTGCTCCCTGTTGTCCCC
AGTCCCGCAGAAGGCTGCCGGGGCGCAGCACCTCGTGGGGGGGTCCCTCC
TGCCGCACCTCTCCCCCTCCAGCACACCACCCGCTCTGCCCGCGCCGC
CAGGGCTGCCCGCCCGCTCACCATCAGCACTGCAGCCCCGACCCTTTGG
CTGCGAGGATCTCCTGCTCCACCTGTGGGAGGAGAAACGGTCAGGGGGAT
GTCTCAGCCACTGCCAGGGACCGAGGGACACCAGGAGTGGAGATAAGGG
GACACCAAGAACAGGGGACCATGGGGACCTAGGCGTGCAATCTGAGGGAAC
ACAGGGCTCAGGGGGATGTGGGGACACGGGGACGTGGGGGACATCCCCACC
TGCTGCTGGCTCTCAGTGTCCAGGGCGCTGGTGTGCTCGTTCAGTATGAG
GATGCGGGGGTCCCGCAGCAGCGCACGGGCAATGGCCACCGCTGCCGCT
GCCCCCGGAGAGCTGTCTCCCACTCGCCACCTCTGCGGGGACAGCG
GGGTGAGGCTGGGAGGGGACCATGGAGGGGACCCGGAACAGAGAGGGGAC
AGCTTACCTGTGTGCTAGCCTTGGGGCAGGCGAGTGATGAAGTCGTGAGC
ACCCACCTGGCGGGCGGCCGCTGTACCTGCGCCCGGCTGCAGCCCCCA
ACCCATAGGAGATGTTGGCGTGGAGTGAGCGGGCAAAGAGCAGCGGCTCC
TGGGGGACGACGGCCACCTGCAGTGGGGGGGACAGCTGGGGACATGGGCA
CGTGGCAGTGGAGGCGGTGGGGATGGCTGGGGATGTGTGAGGGACATCTG
GGACATGTTGGGATGTTGGGGACACAAGGATGGTGGGGACTTGGCCG
GGACACTGCAGGGGACAGTGGTGGTATATGGCAGAGACATCAGGGTGTGT
GGTGTGCTGGGGACCCAGATATCTGGTGAATGAGGATGGCTGCT
GATATGCAAGAACACAGGGGGACAACCAGGAGCCATGGGGACATGTGGCT
GCTCACCTGGCGGCACAGGTAGGAGTGTGGTAGGCGGGGAGGGGGTGGC
CATCCAGCAGCAGGCGTCCGGCCGTGGGCTGGTGCAGGCGGGACACGAGG
GCCACCAGAGTGCTCTTCCCTGCGCCCGGGGGTCCCAGCAGTGCCAGCAC
CTCCCCGGGGCGCAGCTCCAGTGATACGCCCTGGGGACACGGATGTACAC
CCCATGGTGTCCCTGTACCCACACCTCCATAGTCCCACGGCCTTCCTGCT
GTGTCCTTCTGTCCCTGTTCCCCCCCCGACCCTTTGTTTTGTTCCACACC
TTTGTACCATCTACATCCCATGCCCTCCCCACCATGTCTCTGTGTC
ACTCATATGTCCCAGTGCCACAAACCCACCATGTGCCATGTCCCCGCGC
CCTCAATACCATGATCTCATGTCCCTCTCCAGTGTCCCCATAACACCCCC
TCCATGTATCTCCTTGTTCATATCCTCACACCCTGTCCATCACATCCAT
GTCCCTGAATCACCCCACTGTGCTCCACACCTGTACGCTGTCCCACCA
CACCCCGTGTCCCTGGCTGTGTCTCTGTGCCACCTTGAGGACGGGTT
CCTGGCGCCAGGGTAGGAGAACCAGACATCCTCCAGCTGAAGGTGGCCC
TGCAGGTGAGCGGGTGCCATTGTCCCTGAGGGTGCGACCTGTGGCTCCCG
GTCCAGGAACCTCAAAGATCTTCTCCGAAGAGCCCACAGCCTTCATCAGTG
TGGGGAAGTAGTCGAGCAGGACCTGGGGACAGCGTAGTGACGTGGCCAGG
AGGGCAAGGGCATGTGGCATGGTGACATGGGGACGACAGGACACAGTGG
GATTGGCATAGGGACAGGACGGGGTAGGTATGGGGACAGAGGAGGTGACA
CAGGGATGTGGCAGGGGTACAGAGGGTTAGACGTGGGGACACAGGGACA
TGGATGGGCATGGGGACACACTGCGATTGGCCTGGAGACACAGCAGGGAG
GATGTAGGAACACAGGGACACTGTGACACACTGTGACTGTGGCAGGGAGA
CATGGGGACATGCGGATATGGGCATCCTCTCAGGCTCACCTCCAGGACAT
CAGTGAAGTGTATCTGGTAGAGGAGGAAGGTGACGAGGTCCCCAGTGCTG
ACGGTCCCCCGCGCCACCAGCTGCCCCCATAGTAGAGGATCCCCATCTT
CAGGGCCAGGGCTGAGAAGTGAAGGATGCCATGAGGTGACAGGGGACACC
TCCCCCTGGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT

FIGURE 10

SUITE 54

GTCAATGTCCCCCTGTCCCAGTGTCCCACAGCTGTGCTATCTCTGTGCT
 ATGTCCCCTCATGCCATCATGTTCCCAGTGTCTGGAGCCCCCATGCCGC
 CCCATTCCCACGTCACTGCATCCTCCTGCCCCAGAGCCCCCTGAACGTGTG
 TGCCTGTACATCCTGATGTCCCCATGCCATCAAATCTATGTCCCACAGT
 CCCCATGCCATCATATTCCCATGACCTGCCATCTCCACACCATTATGGCC
 TCCAGCCCCGTTGGCATTCTGTCCCCATCTCCTGACATCTCAATTACATCA
 CGTCTCCACATCTCCCAGCCCTATCCCACCATGTCCCCATGCCCCCAGT
 CCTATCCTATCATGTCCCCACATCCCCAGCCCCATCCCATCACGTCCCC
 ATGCTATTGAGCCACCCCATCCCATAACACCACTGGTCCAGAGAGAGGCAG
 TGTAGAGGGCCACATCCTTCTTCTCCAGGCGGTGGCTCTGCTGCAGGCGC
 TGCCGGTAGTGTGCAGCCGCCCATCCTCATTGGCAAAGCTTCGCACAGT
 GGCCATGGCCCTGGAAGGTCTCCACTGCCACCTCGCTGGCCCGGGCCTGCG
 CTTCTGCATCTGTGGTGCCAGGGCCTGGGGACAGCAGTGTCAATTGCAGG
 GCGGGTGGGAAGGGAATGGGGGCTGGGGAGGGGACAGTGCATGTATGGAG
 GGAACAAAGAACACGTGGAGGGGATGGAGGGGACATAAAGGGGACGGTGG
 GCATGTGGAGGTGACAGTGGGGGCATGCAGAGAACAGAACCCATGTGGAA
 GGCATAATGGTCACACATAGGGAATAATGAGCACATGCCACGGATGGTGG
 ACGCATGGAAGGGGCATGGGGCACGTGGAGGGGACAGCGGTACACAGAGG
 GGACAACAGCAGGAGGATGGTGGGTACATGGAGGGGACAGTGGGCACATG
 GATAGGGCAATGAGTACATGGTGGTGACACTGGGCACATGGAGGGGACCA
 GAGGCACATGGAGGGGACCAAGGCACATGGAGGTGCAGACAGCAGCCCA
 TACCTGCCGGAAGTGCCCCACAGCCCTGGGCAGTGCCAGCAGCAGTGGCA
 GCGCCAGCGCGGTGAGCAGCGCCATGCGCGGGGACAGCCAGGCCATGGTG
 GCGAAGAGGCGAGAGGCCACGTGCCAGATACCACAGCAGGAGGCTCAGCGC
 CTCACCCAGCGCCTCGCGCACGTCTCCGCATCCCGCGTCACCCGCATGG
 CCACATCCCCTGCCGGGTGACAGCGCCGTCACTGCCACCCCTGTCCCTTA
 TCCCCGTGTCCCCTCCCAGCCCGGTGCCCTCACCCGCCCCGTGCGCGCG
 CAGCTCGGTGATGCTCTGCCGAGGAGCGCGCGGAAGACGCGCGCTGGA
 GCGGGCTCTGCGTGCGGGTCACTGTCCCCACGAAGGTACATCACACACC
 AGCTCAGTGACAGCGCTGTGACGGCAGTGGGGTGTGAGGGGGTCCCGCGC
 TGCCCCCTGCCCGCACCGCGTCCCCCTATGCCAGTACCTGCTGAGGCCC
 AGCAGCACCATGGGCAGGATGGCTGCCAGCTCATCCTCGCGGGCCACCCA
 GTCGCTGGCTCGCCCCATGTAGTAGGGCACGGCCATCTCGCTGTACCA
 CAGCAGGGTCAGGGCATGGGGGACCCCCCGAGGAATGGTGCCCCAGGAG
 TTCTGTGCTGCACCCCACTTTGGTGCTGCACCCCCAAAGCTCAGAGGTG
 AACCTCCGAAGCTCATTGTTGCCCTCCAGTTGGCTGCAGCCCCCACCCC
 ACCGCAAACCATCTTCATTCCCATCCCATCCCTACCCACCCCAACCCATT
 GACCCCAATTCTATCTCCATCCCATCCCTACCCACCCCAACCCATT
 CTATTCCATCCCATTTCTACCCCAACCCCAATCCAACCCAGTCCCATCT
 CCATCCACACTCCATCACATTCCCATTCCACCCCTGTCTTCAATCCCAT
 TCCATCTCCATCCCAACTCAGCCCCAGTTCCCATTTCTCTCCCATCCC
 CACCCCATCCTACCCAGTCCCAATCCCAGTTCCAAACCCACATCATTACC
 ATTCCATCCCAACCCATTCCCAGTGCCAGCCTATACCCATCCTTACCC
 CCACCCCAATCCCATCCCATTTCCCATCCCATTCCACGGCTACTTCCAT
 CCCCATCCCATTCATCCGGTTCCCAATCCCATCCCATCCCTACCCCTT
 ATCCCCAGCCCCACCACAACCCCATCCTCATTCAAATCCCAACCCCACTC
 CGATCCCACTCCCACTCCCCCGCCCCGTACCCAGCGCCGAAGCCGCCAT
 CAGACCCATCACCGCGGCGCAGCGCCGCGCTCCGGGCTCAGCGAGAGGA
 GGAGGCGGCGACCCGCGCCCATCTTCCCATCGCGGCCCCGATCCCCCTC
 CGGCCGATCCCAATGCCCGCAGCGGCGGAGCTTCTCGGAACGAGAG
 CGTCTCTCATTGGCTGAGGCGGTGCAGCAGCGACGCTGCTCATTGGTGA
 GATGGTTTTCGCGTCATCAGTTGCCAGGCAGATCGGAACACTGCAGTTGG
 AGAGGGGCGGGTGATTGAAAGTGAAGTAACGGCGGAGCGGGAAGGAGAT
 GGAGAGCGGCGCGGTGAGGGGCTGGAGGGGGCTGGAGGGGGATGGACTG
 GTAGGGGCTGGTGGGCTCTG

FIGURE 10

SUIITE 55

204743160 "014402

GTGGGGACTGTCGAGGGGCTGGAGGGATCTGATGGGGACTGGAGGGGTTT
GGTGATCGCTGTTGTGTGCTCCAGGCTGGGCTGTGGGGAGCCGACTGGA
AGTGGGGGCGCTTCTAAAAGCACTGCTGTGTGTTCCAGGTGCTGAGGGGA
GCTGAGGACCTGCACCAGGAGCACCCCGGGGAGCCACCTGGTCCAGCTG
TGCACCAGAAGCTCTGGGGATCCCCACCCACAGCCATGGCGATGCCGCC
CTACATTCTGCGCCTGTCTATGCACGCTGCTCCTGGCCGACCTGGCCCTCA
TGTTGGCCCTGGCCCACTTCTTCCCAGCACTGGCCCACTTGGGCTGGGTG
GGCTCCTGGCTGGAGGCCGGGCTGCGCCTCCTGGTGTGGGGGGGGCGG
GCAGCTGCTGGCCCCCAGGGGACCCCGTGGGGCTGCACTGCTGCTGAGCC
TGGGCCCCGCCATCTTCTGACCCTACGGGGCTATGTAGGTCTGCCTGGA
GCTGCCCCGGTGCTGCTGGCCATGGCAACGCCGCTCCTGGCTGGTGTGAC
CCACGGGACAGCTGTGGTGGCATTGCTCACCTGGAGCCTCCTGGTCCCCA
CTGTGGCCACTGGGGCAAAGGAGGCAGAGGCCTGGGTGCCCTGAGGCGG
CTGTGGCCCTCGCCTGGCCGAGTGGCCCTTCTTGGCTGTGCCTTCTCT
CTTCTCGCATTGGCTGCACCTGGGTGAGACCTCAGTGCCTACTGCACCG
GGAGGGCTCTGGATGTCTCTCCGCCAGGGGACGGCCTCGCCGCTTACC
GCTGCTGTGCGCCTCATGTGCTGGCCTCTGCCAGCAGGTAGGGACCCCA
CATCCCTCCACAAAACCCCATCCACCTCTGGTGGTCTGTGGTGGTTTG
GGGGTCTCTGTCCATATCTGGGGGTCTATCTGATGGGTCTGGGCACTCCA
CTGACCCCTTGTGATTGTCTGAAGGGTCTGGGGCTCTCCATTGACCCCTG
ATGGGTTTTGGAGTCGCCCCCCCCAATTCTTCCCAGCTCGCTGTTTGCCG
GCTGCCGCGGTGGCCTCTTACCTTCATCAGTTCCGCTTCATCTTGCGC
ACCCGCGACCACTCTTCTCCAGCCTGGTGTACCGGGACCTCGCCTTCTT
CCAGAAGACCACAGCAGGTACAGACTGGGGGCACTTTTGTCCCTGTCCCC
ACACCATACCCCACTCACCCTACTCACTCCACAGCTGAGTTGGCCTC
CCGGCTGACCACCGATGTGACGCTGGCAAGCAAGTGTGGCACTCAATA
TCAACGTCTGCTGAGGAACCTGGGGCAGGTGCTGGGGCTCTGCGCCTT
ATGCTGGGGCTGTCCCCGCGCTGACAATGCTGGCACTGCTTGAAGTGCC
GCTCGCCGTCAACGCACGGAAAGTCTATGACACCCGGCACCAGGTGATAG
CAGGGATGGGATGGTAGGGTTGGGGTGACAGGGATGGAGGCAATGGCAAT
GGGATGGGAACAGTGGGAGTGGGGATAGTGAGGTGGGGATTGTGGGGTCA
GGGTGGCAGGGATGAGGGCAGCTGCAATGGGATGGGAACAGTGGGAATGG
GGAGAGCAGGATGGGGATCATGGGTCCAACACAGCAAGGATGAGAGGATG
GAGAAGAGTGGAGCAGGAATGGAAGTGGGATGGCGAGTACTTGGCCATCC
CATGGGTGCTGACACCCACTGTCCCCCAGATGCTGCAGCGGGCCGTGC
TGGATGCAGCAGCCGACACCGGAGCGGCACTGCAGGAGTCCATCTCTTCC
ATTGAGATGGTACGGGTCTTCAATGGCGAGGAGGAGGAGCACCCTA
CAGCCAGGTGCTGGACAGGACCTACGGCTGCGGGACCAAGCGGACACAG
AGAGGGCCATTTTTCTCCTCATCCAGCGGTGAGGCTGACACGAGGGGAC
ACCCTGGTGTCTGGGTGGGATCGGGACATCCCCGCTGAGCCCCATCCCCA
CAGGTGCTGCAGTTGGCCGTGCAGGCACTGGTGTGTACTGTGGGCACCA
GCAGCTCCACGAGGGGACCCCTCACTGCCGGCGGCCTCGTTGCCTTCATCC
TCTACCAGACTAAAGCTGGCAGCTGCGTGCAGGTGAGGTGAGGCAGTGCG
TCCTCTGCCACCGGATCCCCATGACTGTGGCCACATCCCCGTGTCCCCAC
CCTGGGTGCTGTGCCTGGGGGTACATCCCCATGTCCCTATCCTGGGTGC
TGTGCCATGCAGGCACTGGCGTACTCCTATGGTGACCTTCTGAGCAATGC
AGCGGCGCCTGCAAGGTCTTTGATTACCTGAACTGGGAGCGAGCTGTGG
GTGCTGGTGGCACCTACGTGCCACCAGACTGCGAGGCCACGTACCTTC
CATCGGGTGTCTTTCGCTATCCCACTCGCCCTGAGCGCCTCGTCTGCA
AGATGTACCTTCGAGCTGCGCCCCGGTGAGGTGACGGCGTTGGCGGGGC
TGAATGGCAGCGGGAAGAGCACCTGCGTGGCACTGCTGGAGAGATTCTAT
GAACCTGGGGCCGGGAAGTGTGCTGCTGGACGGGGTGCCGCTGCGGGACTA
CGAGCACCGCTACCTGCACCGCCAGGTGAGGGGGTGGGGGGAGATGTGGC
TGCACTGAGCAGTGTGGGGCTGAGCCTCTGCCCTGGGGCAGGTGGCACT
GGTGGGGCAGGAACCCGCTCCTCTCTGGCTCCATTGGGGATAACATTG

FIGURE 10
SUIITE 56

FEUILLE DE REMPLACEMENT (REGLE 26)

TTATTTTACAGGAAGCAGTGCAGGCAAAGCATAACAGTCAGCATAGGAAG
CAGAATGAGATGTGAGAGGTCAAGAGGATGGGGCTGTGCCCATCACTGAC
CCCAAATGCTGGTGTGGACTGTTGGCTCCTTGCAAGGCTGAAGCACAGCAC
TCAGATGGCGGGGTGCTCCCTGTGGGGATGAGAAGGGATGATGTCCAC
GGGGCCAGGTTACATCCCAACCTCTGCTCCGGGGGTCTCCCCGCAGAG
CCCCAGCATAGCCCAGCCCTGGGCAGTACCACAGAGAGCACACGGACCCC
TCCCAACCCACAGCCCCACACCGCACCTGTGCTCGAGCTGCTGGATCCAC
CTTCCCTGTCTATAAGGGAGGAACAGCTGTGAGATCCAGCCCTGTGCTCC
CTATGGGGCGTGGAGCAGCCCCAAGCACCAGGAATGGGTACCCTGGGAGA
AAGTGACAGACCCCAGGACCGCCCTGGCACTGGGCAGAGGTGGGGGACAGC
GCTGCCCTCATCACTACCGGGCGGATGTTGTAGCCCTTCCCCTTCTTC
CCTGCAGACAGAGAGATGCTTCAGCACAGGTTACATCACAGCCAGGCTG
AGGGCTCCCAAGGGGGACCCAGATCCCCCACTGCCCGCCTGCACCCCTC
TGCTTTTACCTGCATGGCGTCTGTAGATGATGAATCCAACACCAACCATG
ATGGCAATGGCCACAATGGCGACGGCCACCCCGCCACGATGGGCACCAG
GTTGGGCTGTGGCGGCTCTGGGGGAGAGCGGGGCCATCAGCAGGGAAGG
GGCAGCCCCGAGCCCCCAGCCCCACATCCCTCACTCACCCACGAGTA
GAGGCCGGGCTGGGGCAGGCTGGCGTGCTCCACGCGGCACTGGTACTTGT
CCCCGTCCCCGGCTGCGCATCGATGGTGACCCAGGTGTGGTAGGTGCCG
TCGCCGTTGGGCACGATGCCCCCGAGTGGGCGTCTGGCCCCGCACCGC
GCCGTCTTCAGCCAGCTGACAACGATGGGCCGCGGGTAGAAGCCGTGAG
CGCGGCAGGACAAGGTGAGGATCCCGTCGGCCTCCTTCCCCCACTCGC
ACCTCGGGCCGCTCTGCGGGCGGGCGGCAGTGAGGGCCGGGCTGAGCTCC
CCACGCTGAGCCCCCGCCCCACGTCACAGCCCCACACTGCAGCCGCTCCCC
CCCCACCCCGCTCACCTCTCCTGCCAGCTCAGCCTTCCCCTATTCCAC
GTATCTCCGCAGCCACTCCACGCAGGTTTCTTCCAGGTAATTCTTCCACC
TCTCAGGTTCACTCTCTTCTCCATTTCTTCTTGGTGGGAAGTGCCTCT
GGAAGTCCCGCAGTGAACGTCATCGTGCCCTTGTGCAAGGCAGTGAAGT
TCTCCCATCGTAGGCCATCTGATAATACCCCGGATGGGGCCGCCCTCGA
GGATGTACAGCCGTACATCCACTGCACCGTGTGAGACCTGAAACACAG
CCGGGCAGGGGGTGGGGGGCCCTCCGGCTCCGGGGCTCCCACTGCAGCG
GGGATGGGTTGGGGTCCCCCGGGACGCGGCACCCGAGCGCGGTGGGGC
TGTGACGGGCAGCCCCGGGACGGTGCCGCGGGCAGGGCCAGGCCTGGGG
GGTGTGGGCGGCACTGCGCCATGGAGCCCATCCACACCCACGGAGCCGC
GGCCCCGGCCGTGCTCACCGCCGGTCTGGTGTAGCGCCGCTGCAGTATG
CCCAGGTTCTCGCGGTCAATCTGCTCATTGCCCTGTCCGATCTGCGTCTG
TCCATCCCAGTACTGCTGGTCCGCCTTGGCCGCTATCCACTCGGTGCGGG
GCACGTACCTCCGCGCGGTGCTGTTGTAGTGACGAAGAGTTCCCCGTCC
ACGTACCCACAGTCACGAACCACGGCTGCCCGGGGCGGGATCCGTGAT
CGCCGTTTGGATGTACCGCAGGGTATGGAGCTCTGCGGGGACGGAGCACA
GCGGGGCCGTGAGCCGCGGGTGTGGGTCTGAGGATCCACGGACACAGCC
CCGGGGTGGGGTTACGGGGCGGGGAGGGGTCCCGGTCCGGCCGCACTCACC
GGCCGCCGCCCGCACACGGCGGCGAGCAGCCCCAGGGCCAGCGCCC
CGCACGGCCCCATCGCCTCGCACCGCTGCACTCTCAAGTCCGCCGCACCC
AGCGCAGCTCCTTCACTTTTCTTACTCCTCCGCCCCGCCCCGCGCCCC
ATTGGCTCCTCCGTGCGCTCTCCGCCAATGGTAGTTGGAGTTTTATGTGA
CGTCATCGGGCGCCAGGCAGAATGCGCTCCCTCAGGTTGTGAAGCGAAAG
CGAAAGCGCGGAGCGGGGGAGGGGATGGGCGCGGTGTGGGAACCCCCGGC
CCTTCGAGCACGGGGGGGACCCGGGCTGTGTTGCGACGGGGCCGCGTCC
TTACCCCGGGGGAGGGGGCGAGGGTCTCTGCCGGGAGGACGGGGGGCGT
GAGAAGAGGAGGAGTCACTTCTCCATTCCAGTCAAGGAAGTGTGGGGG
GGGGTACATCCATAGGGTTAGAGGCTCCGTGTCCGGGGGGGAGGGGGT
GTGACAGTGGTGTCCCCAGGGCTTCTTGGGATCAGTGCCATTTCCCC
ACAGCGCCGCCCCACACCGCTTCCCCACATCCACGTGGTCCATCTGAGGT
CGATGCCCTCAGGGTCTGCAGGTGGACCCCAATGTCCACCCCCCAAGTTA

FIGURE 10
SUIITE 58

10043160 "011402

ATGATTGACCCCAACCCCGCTGTCCCTGCGCCACTGCTCCCATCTGCCCC
ACACTGCCGGAGCCATGGGGCCTCACTGGGCCTTCAGCCTCTTCCTCCTC
CTCTTCCTCACTCCCTTAATGAGGGCCAGCTCCCAGGACCCTGAGTATGG
GGCTATGGGGTGTGTTGTGGGGTAGCTATGGGACTATGAATGTTCTGCAGT
GCCTATGAGGGACTATGGGGCACTGGTGGGGCTGGGGGCTGCTATAGGAT
TGGGGTGTGATGGAGTCTGGGGGGACTAAGGGAGATTTCTGTGTGGTTGG
GTGGGGTTATGGGGCCAGAGCTGGGGGGATTCTCTATGGACCTAAGGGGTG
TCTGGATGCTTATGGGATCTGGGAGGGCTTATGGAGCAGTTATGGGGCTG
GTGGCTCAAGCAGTGTTCCTCAGGTTGGTGCTGGTGGCCCCCGGCGCG
TGGCCTTGGGGACCCCATGGGGCTGTTGCTGGCAGCTGTGGGGCCGGTG
ACCGGGACGGTGA CTGCATGGGCTGAGGGGGACCGTGGGGCTGGGCCCTG
CACCTCCCACTCCATTTGCCCTCACACCCCAACAACCTTCAACCAGC
TCCTACAAATTGAGGTATGGGGACACCGGGGGATATGGGGACACTGGGGG
ATGTCCTCTGGGGTGAGGGGGTTGGGGACACCCCTGTGGCACACAGGGAT
GTGTGCACCTTGGGTCCCCCTCCTGCCATGTCAACCATGTCACTCACAT
CTCCTTCCCCAGAGTTCCCCCATGTCCCCATAACCCAAACACCTCCTGC
TGTGTTCCCATGTCCCTTATAGGTCAACCCAGTGCAGGCAGAGCGCTGTG
GGGCGCTGTGGGGTCGGGGGTGCTCCTGGAGGCCACAGCTCCCATCTG
CCCCCCCCCAGTACCAGGAGTCTGAGTGTGGCCCTGGGGGGGCGCGGGG
TCACCTCATTGTGCAGACAGACAAACCTCTCTACGCCCCCGACAGACTG
GTGAGTGCTCCACGTCCACCCTAAAGCCATCCCTCATCTGCCACAGTTC
TCCCCCAGTGCCCCAAATGCTCCAATTCCCCTAAATCAACCCCAAAT
CTCCCCAAAGCCCCCTCAAATCTACCATGAATCCCCAAATCCACCCATT
TTCACCTACATTCAACCATTTACCCCAAATTCACCCCAAGCACACCCCA
AATACCCCTGGTCAACCCAAAGTCCCCCAAATCCCCTTCAAATTCCTAAA
TCCATAACCCCATCTGTCCCCATGTGTCCCTTTGTCCCCAGTGCGTTTC
CGGGTCTTCTCCATGGACCCCGACCTACAGCCGAACCCCGAACCTGTCTT
GGTCACCATCACGGTATGGGCCCTATAGGGCTGGGGCTGTGGGTGACCCT
GTGGGGTTTGGGTGACCCTACAAGGCTGTGTACCCCATGTACCCCAAG
ACCCGTTGGGTGCACGAGTGCAGGAGGTGCAGCGGGTGGCCCTGGACACG
GTGCTGAGCGACAGCTGGTGCTGCCTGACATCGCCCTGTGAGTGGGGCT
ATAGGGGGCTACAGAGGGCTGTGGGGTGCGACAGGGGGCTATGGGGACTG
GGGACTATGGGGATTTGGGGCTACAGGGGGCTGCAGGCGGGCTAGAGTAGT
GGGGGGGATTATAGGGTTACTGGGGCATTACAGTGGCCATAGAAGCTATA
GAGGGCTGTGGAGAACTATAGGATACTTAGGGGGCCATAGGGGTCTACAG
GGGTTATAGGTGAGCATGGGGAAACATAAGGGCCATAGCGACTCCGGAGG
GCTGTAGCACACCATAGGGGGCCATAAGGGCCCTGGAGGGCTCTAGAGGAC
CACAGAGGTGTATGGGAGGGGGCTATAGGGGACTATAGGGTATAT

FIGURE 10
SUITE 59

10043460.01402

97/110
E52FOR.txt

ACATGGGAACACATGAGGACAGGGAGAACTGCAGGGACACAGGGACACT
TGGGGGATAGGGGGATGGTAGTGATGCATGGGGGGGGGCACATGGGGATG
TGTTGGGGCACACTGGGATGTGTGGGGATATGGGGACACATGGGGAATAT
GGGGATGAGTGGGGACATATGGTTATTATAGGGATGTATGGAGACATTGG
GACACATGCAGAGGAGGGGACAAATGGGGACACACTGGGGGACAGATAGG
GACATGGGGACACCCAGGGAGGGACACCCCAAGTTCCCCCTTACCGGCGG
CAGTGATGGTTCCTTCTGTGCCCATCCCCCCTGCAGCAGCGCAGTGACA
CCGTACTGCGGGGTCCCCACCGCCGCCACCCACCACTGCCCCCCCCGCGGT
TGGGGGGCTGCGGGCGTTCGGGGTGCAGAGGGCGGCTCCATGGGTCAGAGC
CGGTCTGGGGGTTTCGTGGGGTTCAGTTCGCAGCTGGGGGGAGTCCGGGGG
GGGACCCCGAGTGGGGTCAGAGTCCCCAGGGGTCTGCGAGGGAGAGAGG
AGTGAGAGGGATGAAGGGGTCTGAGGGCATGGGGTTGGGAGGGGTGTGGG
GCGTAATGGGGTCATTTTGGGGTTAATGGGGACACTGGGGACAGTTTGGG
AGCTATTGGGGCTAATGGGGTCTCTGGGGGACATGGAGGGGACATTGGGG
ACATTTGGGGTGTAAAT

FIGURE 10
SUITE 60

2044360 "01442"

TGGCTGATGGGCTGTGTCTATGAGCGCAAAACACCACAATGGGCAGAAA
AACCTTCCTCCAGAGGACCAACCCCATCTCTATGGCTTCTTTGCACCTGG
CCTTGCCCAAAATTGGGTATTTTTGAGAAAAAATGGGCCATTTCTCTG
CTGGTTGTCCAAGCAGCAAGAGATGCTGGCATGAGTCTCACCAAGCCAAG
AGGTCTGTGGGACCAAGAGAACTCTTTTCTCTCCCATTAATGATGAGTAA
CTCCACCTTTGGGCACTCTTAAGGTGAAAATCCTCAAAATCTGCAATTTT
GAAGGCGCAGCTCCACATTTCTCATCCCCCTTTGTTCTGTCCATGGCAGT
GCAGGCATTCAGCCCCATCCCCAGCCCTGTGCTCAGTGTCCCTTCGACT
GGATTGGATTGAGAGGAAAATGCTACTACTTTTCAGAGGATGAGAGCAAT
TGGACGAGCAGCCAGAACAACCTGCTCTGCTCTTGGTGCTTCCTTGGCTGT
GTTTGACAGCGCTGAGGACTTGGTGAGGGGGACACAAAAGAGCCACCAAT
GTATTTTGTCCGCTTGAGGGCCCCCTTGGCTGCTCTTTCAGTGTTTCCTTT
CTGATTTTGGGGTGAGGAGGTGGATAATGGTTGTCTGAGGGTAGGTTGG
GTCTACTCCTCAAAATCTTCAAGGGATTTAAGGGAAAAAAGATGTTT
TTTCTATGAAGTAACACGCTGGGTTAGAGACTGTGAGCTTTGGTGATGG
ATTGGGCAGTTTCAAGCACTGAGATTATTGGTTGAAAGGGTTCTGCAGGC
AGTGGCATGCAGGAAATGTCCAGAGCCCCATGATCTGTTCCCTCTCCTC
TTTTCCAGAGCTTCACAATGAGACACAAAGGCAGCTCCCCCACTGGGTT
GGCCTCTCCCGGGAAGGCAAAGAGCATCCATGGGAATGGGTGAACCGCTC
TCCTTGTCTCACCTGTGAGTTCCCATCCTTGTCTTGGAGGCTGCAGTT
CTCCAGCCCCAAAATGTGGATTTCTGGACCTCGGGAGCATTCTGGAGGT
GGCTTATGGGGTGAGGAGATGTGGGGAAGGCACTTCGCACCGCTTTGGGT
CATAGAAGTTCATTGAGAGGCAGAAGTGGCGCAGGAAAAGAGATTCTTA
TTAATCAATTATTTTGTCTGTTTGATTTCACCACTGTGATTTCTCTTC
CCCCCCCCCAMAACCTGGGGTCTGCCTGTCCGTCTGTCTGTCCATCCA
GGTTCAGGTGCAAGGCGATGGTCTCTGTGCATACCTGGGGGATGCCGGG
CTCAGCTCCTCCCACTGCAGCACGCGGAGGAATTGGGTTTGCACCAAACC
CGCGTTGCAAAAACCGAGGAAGAACTTCTGCATCAGCACCTGAGCGGCTC
CCGGACCCGAACACGCGATGCAAGAGGAGGAACCCAAAGCAAAAGAGCTC
CGCTTTCAGCTGTGCTCAGTAGCAACAGGAGGGCGGTGCGCTCCTCCAGC
CCAGGTCCGACAGTGCCGCTATGGGGCTGCGCGGACCGAAGCAAATCCC
AGGCGGAGCTTCGGCTCCAAATTACATTTTTTTGCACCGTCTGACTCCTA
ATGACCGCTAAAATCCCAATTTTGGGGGCTATCCGTGCGCTGCTTGCAAC
GACCTTCACCCCTGCGCGATGCAGCAGCAGGTTTGGGGGGCGGACGGTGG
GAAAATATCCATTTTTTACCAGTTTTTCTCAAAGGGAAATACTGGGAAA
GCAATCAGCCCAAAGGACCCTGAAATCGATGAATAAATCGGCAAATTATT
TATGTTTTCGTGTTTTCCCTTCTGTGTCACTGCAGTGCCTTCTCCATGAAT
TCACTTTTAACGGTGTTTTGTACAGGAAACACTTCTTCGACTCTCTCCA
CCACTCCTATATATTCAACAGACCAATTCTTCTGGTGATTTTATGCAAA
AACAAAAGAGTATATTTGGTTAAAGAACCCAAACCACCTTCTTGTACTGA
AGGGAATAGAAGAGCACAGACCGCCCGCTCCCCTCCCCTGCTGCCGCACA
ACAGACGGTCCCCGAGGATGTGCAGACAACGCGACGCCGTCTGA

FIGURE 10
SUITE 61

10043450 01402

99/110

E6G2N15.txt

TAGNAACTAGNGGATCCCCGGGCTGCAGCTATGGGGGAGTGGGTGCACT
CCTTGGCCATGGCTTTGGGTCCCGTTACTCTGGAGGAATTTCCACAGCTG
CCCCAGGAATCTTGACATAAAAGTGCACAGATCGATCAGAGATGTCATG
TTCCTGACAGAAGAAATCCTGTCTCTTCTGATGTTCTCTGTGAAGAGCAT
TGCCACGAGGGAGCTACCAGCAGGGCAAGCAGAGAAATTGAAGAAAACGA
AAGATGGGTCGAGGTACGGGATTGGGCAGGTTTCACTTTCTTTAGCAATG
AGACGTGTCAAGCTGGCAGCTTCCCTGGGAGCCTCTCTGGTGTGGATCTC
CGGTGGCCCTAAACCTGGTTCAGGCACTGATCAAGGAGACATTACCCGTC
TTGGTTCATCTCGGCTCACGG

FIGURE 10

SUITE 62

2044160.03440

100/110
F12FOR.txt

CGATGGTCCTCCAATGACCTCCATGGTCATCCAGTGCTCATCCCGCGGTA
TGGCCATGGTAACCCCATGTTACCCTGTGGTCTCACCCCAATGATGCCG
TGGTTACCTTTTCGTTACCCTATTCTCATCCCATATCCCCCTTTCTGTC
CCTCTGCCCCTTCATGATCCCCCTCATGGTTAACAGACGTTTCCCTCTGCG
ATCAGGTGATGTTACAGCACAAATTCCTCCAGGGTTCCCTTTATAGTGACC
TCACCATTACCCAATCATGTCCCCGGTGTCCCTGAAGGGGGCCAGATTC
CTCAGTGGGACCCAGATGTCTTCAGTGGGGCGGGACCTGGCCATTCCCAA
TGTCATCCAGGTGTCCATATGGCATGGGACACAGATGTGCACATGGGATG
GGACCCAGGTGTCCCCACTGTCATCCAGATGCCTCCATGGGTTGGGAAAT
GACCATCCTCGATGTCACCCAGATGCCACATGTGATGGGACGTGGCCAT
CCTTCATGGCATCCCGATGTCCAGCTTGGGATGGGATCCCAATGTCACCC
AATGCAATCGCAGTGTCACCCAGATGTCCACAAGGGATGGCACCCAGATG
TCCCCAGGTGCCACTCATCTGCCTCACCAACCCAGGACTTCCTCCCACTG
CTCCCACTGCTCCCAAGTTTGGCCCCCATTTCTCCC

FIGURE 10

SUITE 63

20470 "094001

101/110

G2M13.txt

GATCTTCAGTGATTTTTCAGTGGTCTTTGGTGGTCTTCAGTGCTCTTCGTT
GGTCTTTGACAAAGATGCAGAGGAGCACCGCTCCCAGACGGACCCCCCGG
GGACCCCATTTGTGCCCATCCCCACTGGGACATGCAGCCATTGACCACAG
CCCTCCGGCTGCGACCACCCAACCTGATTCTTATCCAAAGTCCACTCTTT
GCACACTTACCTCCAATTTAGTGATAAGGATGTGGCGTGGGACCGTCCCA
ATGGCCGCACACAAGTCCAGGTAGATGATATGGGATGACCATGAAGGGAT
CACAGAGAGGAACACGGGGTGACCACGAGGAGCAACGAAGGAAACGCTGA
GTGACCACGGGCAGAAAATGGTGTGACCATTAGGGGACAACGAGAGGGAA
CAGAAGTAGTAAGGAGTGAGAATGGGGTGACAAAGAGGTGACCATGGCAT
AACTTTGATAAGACCATTGGGTGACCGCAGGGTGATGGCCATACCATGGG
GTGAGCACTGGATGACCATGGAGGTCAATTGGAGGACCATCGGGTGGGACG
AGGGCCCGTGGGGACACCCGTGGGGCGGTGGGACGGGGGCAGAGTGTGAGA
AGGAGCCCCCGGGCGCAGAACTCTGCCTGGAGACGGGTGACGCCGCCCGG
CGCCGCCCGCGCTCATTTGGCCCTCCCCGCCCGGGCTCGCGGGT
GCGCGGGGTGCCGGTCCCCCATCGTCCGGCGGCAGCAGCCATGGGGAGC
GGGCGCGTCCCCGGCGGGGGCGGTGCTGGTGGCACTGCTGGCGCTGGG
AGCCCGGCCGGCGCGGCACGCGGCCCTCGGGTGAGCTCGGAGCCGCGG
CGCGGGGACGGCGCTGCGTCCCCCGGAGAAACCCCGGAGCCCTTCTG
GCCGTGCGCAGCGCTCGGGGTGCGGGGGACGGAGGGCGGGGGGGGGCG
GCGGAGCCGTGGGGGGCAGCGGGGCCGGGAGGGGGCGGGGGGGTGTGGCG
GGGGGGCGCTGTGTGCCCTGACCGTGCCCTCTGCCCGCAGCGTTCTTCTT
CTGCGGTGCGATATCCGAGTGCCACTACCTGAACGGCACCGAGCGGGTGA
GGTATCTGCAAAGGTACATCTACAACCGGCAGCAGTTCACGCACTTCGAC
AGCGACGTGGGGAAATTTGTGGCCGATTACCGCTGGGTGAGCCGCAAGC
TGAATACTGGAACAGCAACGCCGAGCTTCTGGAGAACC GAATGAAG
TGGACAGGTTCTGCCGGCACAACACGGGGGTGTGGAGTCTTTCACGGTG
CAGAGGAGCGGTGAGTGCCGCGGGGCGCAGCGCGGACGGACGGGCAGGCG
CCGCGCTCTGGCGGTGCGTCCGCAGCGCTCCCCCGTGCCCCGCAGTGGA
GCCAAGGTGAGGGTCTCGGCGCTGCAGTCGGGCTCCCTGCCCGAAACCG
ACCGTCTGGCGTGCTACGTGACGGGCTTCTACCCGCCGAGATCGAGGTG
AAGTGCTTCTGAACGGGGCGGGAGGAGACGGAGCGCGTGGTGTCCACGGA
CGTGATGCAGAACGGGGACTGGACGTACCAGGTGCTGGTGGTGCTGGAGA
CCGTCCCCGCGGCGCGGGGACAGCTACGTGTGCCGGGTGGAGCACGCCAGC
CTGCGGCAGCCCATCAGCCAGGCGTGGGGTAAGGCCCGGGGCCCTGCC
CGCCGCGGGGGGAGCGGGAGCGCGGCCCGGGCGCTGAGCCGCCGCTTC
GTCCCCGCAGAGCCGCCGGCGGACGCGGGCAGGAGCAAGCTGCTGACGGG
CGTGGGGGGCTTCGTGCTGGGGCTCGTCTTCTGGCGCTGGGGCTCTTCG
TGTTCTGCGCGGTGAGAAAGGTGAGCGCTGGGGAGGGGGGCTGCGCCG
GGGGGGTGGGAGCGGGGGG

FIGURE 10

SUITE 64

20479" 09E400F

GCTCTAGAACTAGTGGATCCCCCGGGCTGCAGGATTACCAGTGTCCCCAA
CTGTTTTTTGCCAATCCAAGCCCTGCAAATGTACAAATATATTAAAGTGGTT
TCCTTAGTTAGACATCTTTATATCTCTCACCAATCATTTAACGTTAACCTT
ACTCTGCTTTTCTTCTGTGAACAGAAAACAAAATCGGAAGCCTCATATACA
GGTGTTCAGAGGAAAATAGTAAGTGGTGATGAAACTTGGAGAACTTGTGA
AGTGAAATATGGGAGCTACTGCCTCTGGAGGGAGGAAAATAAGGAACCAA
TGAAAGATGCCAAGGTGAAGCAAATGAAGGACCAGCTGTTTGTGGCTAGA
GCATACATATCCAGTATTGCTAAAATGCCTTCTCAAAGCAAGTTGACTCG
GGATATGAAACAGAATATCCAAGAGTTTGAGCGTATTCTTAGTGAAAGTT
CTCAAGATGCTGACCTTCCACCACAGTAAGTTCTCTCCAGTTTGGGTTTA
ATCATTTTTGTACTGAAAGTTTAGTTCCTTACTGGAAAAGATTTTGTG
GATTTCTAGTCACATGAATCTCTCCTAGTTTGCCTTCAGTTTGCCGGACA
TCCCGTTTTCTAGTGGTTTTACTTGCTT

FIGURE 10
SUITE 65

20443150-01402

103/110

H4212.txt

TAACCATGAGTGATAAACTGCGGCCAACTTACTTCTGACAACGATCGGA
GGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAAC
TCGCCTTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACG
AGCGTGACACCACGATGCCTGTAGCAATGGCAACAACGTTGCGCAAATA
TTAACTGGCGAACTACTTACTCTAGCTTCCCGGCAACAATTAATAGACTG
GATGGAGGCGGATAAAGTTGCAGGACCACTTCTGCGCTCGGCCCTTCCGG
CTGGCTGGTTTATTGCTGATAAATCTGGAGCCGGTGAGCGTGGGTCTCGC
GGTATCATTGCAGCACTGGGGC

FIGURE 10

SUITE 66

20043160 031402

104/110

H424.txt

TCCCTAGTAACGGCCGCCAGTGTGCTGGAATTCGGCTTAGCGTGGTCGCG
GCCGAGGTACATACCCTGCCCGCAGTGATGTCTCCAAGGTTGATTTAAGC
AACCAGCTCCTCCCTGCCACGGCTCCAGGCTCCACATGCCTGGGTAAAGG
CTGGGTTTGTGTTTTGAGACAGTGTCTTAACTATGGAGCGCTGACTGTTT
TGGAACCTCGCTCTGTAGACCAGTCTGGCCTTGAACTCAGTGATCCCCCTG
ACTCTGTCTCCAGAATGTGGATTCTCCCA

FIGURE 10

SUITE 67

10043130-01409

GGATTCTGACACCCCTCCTCCCCACCCCCAAAGGTGTTCCAGCGCCGCA
 TGGATGGGGGCACCGACTTCTGGAGGGGGTGGGAGGAGTACGTCCATGGC
 TTCGGGAACGTTTCTGGGGAGTTCTGGCTGGGTGAGGACCCCCAAACTTG
 GGAAGATTGAGGTCTGGGGTGGGGGGGGGGAACACCCAGGGCGGAGAGGG
 CTGATGGCTGCAGGACGTGGAGTGGGATCCCTGACGGGGGTGTGGGGTGG
 GGGGTGTGGGGCAGGGGCCCCAGGTGGGTGTGTAGGGTGGGGATGATGAC
 GATGGCTGTGGGATGTGGCGCAGGGAATGCGGCGCTGCACACACTGACAG
 CTTCCGGGGCCACGGAGCTGCGTGTGGACCTCTGGACGCCGTGACACAGC
 GCCTTCGCCCGCTATCGGGATTCGCCGTGAGTGGTCTTGAGGACAATTT
 CCGCCTTACCTCGGGGCCTACAGTGGCACAGCTGGTGTGTGTGGGGCAG
 TGGGAGTCTCTGGGGGATATTAGGGTTAACCTTGACCCATGAGGGGGGT
 TTTGGGGATACCCAGATCAGGGGGGGGGGAATCCTGGGGAGAGTAGGGG
 ATGGTCCCTTTGCCACAGTGAGGGGGCCTTGCCCTTGACAGAGGTCTTTAA
 GATCCTTGACCTGTTGGGATCTCTTGGGGATCTCCAGACTGCAGGGAGCC
 CCGGGGGTTCTTGGGGGGCTCTGCCCCACAGGGTGGTCTCTGTGAGGGTG
 TGGGGGTACCTGGGGGGTCTGCGGCTCATCCTTGGGGCTCTGAATGCTAT
 GTGGGTGTCCTGGAAGGCTCTCTTAGGGGTCCCCATAACCTTGCTGTGG
 GTCCACACAGGGGATGCACTGTCTTACCATGCTGGGAGCCCCCTTCTCCACG
 CGGGACCACGACCCCCGAGGCCGCCCTCGGCCCTGCGCCGTGCGCTACAC
 CGGAGCCTGGTGGTACCGCAACTGCCACTACGCCAACCTCAATGGGCGCT
 ATGGGGTGGCCTACGACCACCGAGCATGGCTATGGGGGTGTAAAGGGGT
 CTGTGGGGATTGTAAAGGGGTCTATGGGGGTATAAAATCAACCCAATGGG
 ACAGGAGGGGGTCACCATGAGGCCATGGGGGTTTTGTGGGGTAAATGTGG
 AGGGTACCCCCCCCCCAAGGTCTTTTAGCCCCATGTCTCTCTGTATG
 AATATGGAGCCCTACAGGAGCTGTGGAAGCTGGAACACAAGCTGGAACAG
 GGAGGGGATACTTTGGGCCCCCCCTGTAAGGCCTATATGTGTCTATAGGGT
 CACTGTAGGTTGTTTAAGGGCATGACCAAGTCCCCCTTCTTTCTGCAGG
 GCATCAACTGGTACCCCTGGAAGGGCTTTGAGTACTCCATCCCCTTCACA
 GAGATGAAGCTGCGACCGCAGCGTGACTGAGAGCACTAGAAAGGTGCTGG
 GTCGCACTGGAGCCTTTATGGGGTCAATAAAGCTGCGAGTAGCCAGTGCT
 GACCCATGTATCCACACACTGGGCTCAGGAGCTATGGGGGTGGGCAGGG
 CGTGAGGCGCACGCGGAACGGGGCACAGCGCAGCACGGTGCCAGCAGTGA
 CCCCTAAGTGGGGCAGAGCCCCATCAGACGGTGGCTCCAGGCGGAATCGC
 TGTAGGATGTGCCCCAAAAACACAAAGAGCTCTGCCCAGCCAGCGCCTC
 CCCACACACGAGCGTGCCCCACAACCAAGGGCAGCAGCGCTCGCCATG
 GAGGCTCCGGTGCAGGAACCGCTCTGTGGGGCAGAACAGAGATCAGAGT
 GGGTGTAGGGGAGGAACCCAGCCTGGGGTTCAAAGCCACATCTATGGG
 GTGGACCCACACATACCGGGCAGGAACCTCATCAGGACGGTCCCAAATCTT
 GGGGTGCTGGTGGCAGCAAAGAGGTTTGGGATAACGATGGATCCCGCAG
 GCACTGGGATTCCCGCAATGCTGGGAAGGGACAGAATGCTGATAGGATGG
 ACTGGGAGAGCCTACAGAGGCCAAGTGGGACATACTGGGACCTGCTGAGC
 TATCCTAGAGCTTACTGGGTGCTTGATGAGTTCTACTGGGACCGACCTAC
 TGGTAGGTCCATGCTGGTCTGTAGTGGTCCACACTATTACAGACTGGTCT
 ACAATGGTTTCATTCTAGTGCAAAATACTGACACGCAGTGGTGCACGGTTC
 TGCTCGCAGACATGTGGCCCGCACTGGTTGGTACTGATCCCCACTGGTCT
 GTATGGCCCCATAACAGCCCGTACTGGTGTTACTGGCTGTACCTGGAGTG
 GCGCCGGGCACAGTGGGGCAGCGCGAGGGGCACGGGGGGTCCGAGGCGGA
 GGGTCTCGGTGACAGTGGCAGGAGCAGTGGCAGTGGCCCCATATCCCT
 GGCTTTGGGGTCCCCCTGGGGCCAGCACCTGGCGCAGCTCTGCACGTAC
 CTGGTCTGCACTGGGACAGGGGACACGTGTCAAAGCACGTCAACCAAGT
 GCCACATCGGGTCACTTGTGGGGTGGCCCTCCCCTGCACGGGGACACAGG
 CAGCAGCGTGACACGGAAGTGACATGAGCGTGACATTTTGGCACTGGCCA
 CAGTGCAGGGGACACCAGGGGCATTATGCACACAGGGTTATGGACATGGA
 TGTGACATGCATATGGGGAAGTGCAGTGGAGCTATGGGAGGGGACAGCCA
 GGACATGGGTGGGGGCGGCCAATGGGACCTGGGACAGTGGAGGTGTGGG

FIGURE 10

SUIITE 68

TGTGACACAGATGTGATGTGGTGTACCTGGGGGTGGTGCAGCAGGAAGG
 CCACAGCCCATAGCAGAGCCACTGCCGTGTTTTGGTGCCACCGATGAAG
 AGATCCACGAGGGCCATGTGCAGGCGGTCCCCCCCCAGCGGCCCATAGG
 GACAGTGGGGTCCCCCCCCAGCAGTGCTCCCAGCACTGTGTCCCTGGGGG
 GAGACGCACAGCCCTGTGGGGACACACGTGTTACCCCTGGGGCCCTGTC
 CCCCCCTGTACCTGTGTCCCCACGTTCCCCACCTGGTGCCATCGGATCT
 GGGACTCCACAAAGGCATCGCGGCGCTCCACCAGGCGCAGCAGCTCCCGC
 AGCCCTGCGTTGGGCAGCACCTGTGGGGCACAGGGACCCCCCAGTGCT
 CCACAGAGCACCCCTGGACCCATAGGGACCCCATATTCCCTCCCAGCCCC
 ATATATAACCCCCCCCCAGGGCGATATAGCCCATCCTTAGTATAGACCCC
 TGCAGCCCCATATGGACCTATACCACCTCCTCTTATGACTATATCCCGCA
 GCCCCACGCCGATCCTATATGCCCTGTAGGGCCCTGTAGGGCTCACCCCTT
 AGTGAAGGCAGCACATCCAGTGCCCGCACACTGGCCCCGGCCCCACACCTC
 CAGCAGTTCACCCACACAGCGCGTGAAGGAGCGCACCTCCGCCTCGGGGG
 GCATCTGTGGGGCACAGGGCTTGGGGTCACCCAGAGAGACTCCTGAGTC
 CCCCCAGAGACTCCTGAACCCAAAGAGGTACCGTGGTCATTTGGATCCCT
 CTAGAGGTGACTGGGTTCACAAAGGGACACCTCAACACTTGTGTCCCTT
 CAGGGGCACCTGGATATCTGGGACTCCAAGTGGCACCTGAGCATTTGGGA
 CCCACCCCTCCTTGGACACCTGGGTCACCCCAAGGACACCTGGGACCCCTT
 CAAGTGGCAGGTGGACATCTGAGCCCCCTGTAGTGGCACTTGAGTCCCCC
 TGCTCCCCCAGGTGACACCCAGACCCTGCAGCCCCCTCGATATCCCCACCA
 GGTCCCCGAAGGCAAGGCGGCAGATGGTGTGAGGTGTGGAACGTGAAC
 GCCTCAAAGAGGTCCACTGGGGCAGCCCCATAAGAGCTCAACTCCTGTGG
 GGTGAGAAATGGGGTCACTGAGCGGGTGCGGGTGCCCCACAAGGGGGGT
 GGGGTGAGTCAAGGGGACGGGCAGCACAGCCCTGGGGCTGATGGGGTCCA
 CCTGGGGTTGGAGGGCCCTGTGTGGGGTGCTCACCTGGCACAGCGCCCA
 GCCCTGCAGCTCCAGGAGGGGGCTCCAGGTGCCTCACAGCTCGCGCCAGTG
 CTCGCCGCGTTGCCCCCCGCTGCCGTGCGCACTCTGGGGATGCATCCCCC
 AGCGCCAGGTCTGCCCCCCCGCGACACCAGGGACGCTGTGGGGTGACA
 CCCATATCACCTGGGCACCCATGTGACCTCCGAGAACCCCTCAGACAGCT
 GTACGGATCCTTGGGGACACATCCAGAATCCCCCAGGCACCCACTGGGAT
 CGCTCCAGCACCCATGGGGACTGTTAGAGATCTCCTCCCCCCCCAAAAAT
 ACAACCAACCCCTTCAAGATCATGGGGACCCCCCAGTACCCCTCCA
 GATACCCAAACAGTGACCTATAGAGACCTCCCTCCACCCAAAAGCCATGGG
 GACCCCTCAGGCCCCCCCCCCCCAGACACCAATTAGTACCCCCCAGAACCCT
 TCAGAAACCTACAAGGACCCACCAGAACCCCTCAGATACCCATAGAGAT
 CTTTACAGACCTCCTCCTGGGACCCTCCCCAGGAGCACAAATCCCAAAGA
 ACCCCCCCTGAAGATTCACAGGGACCCCCCTCTGACTCACCCCAAACCCT
 CATGGGGACCTCCCAACCCCTACAGCCCCCCCCATACCAGGTAACCTGTGG
 GGGCGTCCACGAAGTCCCCCAGCGCCGTGCCAGTGCCCTCACGGATGGC
 TGCTGCAGAGCTCAGCACCACCACCTCTGGGGGGGTGGGGGGGGAGGGGC
 AAAAAATGAGTGAGTTGGAAGGAAGGGACCCCATGGGGACCCAAAAACC
 AGGGAGAGGGGAGAGGTGAGGGGTGCCAGAACGGAGTTGGGGGGGGGGGG
 GGGAGGGATCCCAAATTATTTTTGGGGGGGGGGAGTAGAATGAGAGGAC
 AAATTTGAAGGGGAGCAGAAGGGAATTGGGGGACAGTATGTGGGGGTTC
 TCCATCCTCTCAATGGGTAATTCTGGGGAGCCTGTGAAGTTGAGGGTCT
 AAAGGGGGAAGGCTCAAGGTCCCAAGGAGGGAAGGTTATGGGGAAAAGG
 GGGTAAGTGGTCCCAAGGGGTATCAGGGGGATGGGGGGGGGGGGGGGT
 CATGAAGGTGCCGCCCTACTCACCACACCCCCCAAGCGCAGGCATAAG
 GGGTCCCCGTAGGTCCGGGCAAGGATGTGGAGGTGCCGTGGCCCCCCTGG
 GTGCAGGAGGTGCAGGGCCCCCCCCCACGGTGCTCCCCCCCCGCTGGCTG
 ACCCCCCCAGATCAGAGCCAGGAGAAGTAGCAGCAGAAGTATCGTCACC
 GCCATTGTTCTGTGGGGTGGGGGGCCCCAGCTCTGCCCTATAACACCTT
 ATGAGGAGGAGGTACCCCAAAGCTCCACCCCCCACATCCAAACCCCTC
 CTACCAGAAGAGGGGCATTGGGTTCACTCCCCTAAATTATTGTGTGCC

FIGURE 10

SUITE 69

ACCCCCCTCTTCAAGTCATTATAAACTTTACAGGGGTGTCCTCATAAAAA
TACAGGAGTGTGTCCCCCACAAGTGCTCCCAGAACCATCGGGTGCCCA
TCCCACAAGAAAATCTCTCAAACCTCCCATGTGTGCCCCCAACCAACAA
AGATTCTCTCAAACCTCCCCCCCCCGCCCTCTACCCATATATCCTCCCAAGC
GCTCCCCACCCCTCCGCACACCACCTCCCCAAATCCTCCCCATTACCAT
AATCCCCCCCCACCCAGCAGCAGAACCCCATCACCGCTCTGTGCGTCTGT
GTGTGTGAGTAGGGGACGGGGTGTATTGAGGGGAGGGGGGAGGGGGGA
GAGCGCTCAGAACCCCTCCCCCTGCAGCCCCCGCAGGCGCCGTGCCAGCT
GCAGGTCTTTGGGGTACAGTGTGACGCGGCGCGCATGCAGCGAGCACAGG
TAGGCGTCTTCCAGCAGGTGCACCAGGAATGCCTCCGCCGCTGTGGGAC
CCCGGCGTGGGCGTCCCCACAAAGCAGGGGGGGAGTCAATTCCCACCCCC
AGGCCACCCCCACAAATGCCAATCCTCCAAAATAATCCCTGGAACAACCCC
AAAAAAACCCCTACCCCCAACCCCTCCCCAAAACCATAACCTCAATAA
CTCCACACCTCAAAAACCTCCAACCCCTCCAAAACAACCCCAACCCCGA
AACACCTCACCCCCAAAGACCCCTTCCCAAGCCCCAAAGAGACCCCAAG
CACAAGGGGTACCCCAAATCCACTTCCCCCTTCCCCCAAAAAGCCCTT
TTGGGCACTAGAGAGCTCCCCAGCACCACCCAAAGGGTCCCCACGGTAT
GGGGTACCCTAAAACACCCCCCAACCCCAACCACGGGAATTCCAAAAC
AAAGCTACCCCTCCCCCCCCCCCCAAAAAATAAACCCATAGGGCCCC
CCACCTCCTGTAGGGCCAATAGGGCCATAGCCTGCCACCTGTAGTCCACG
CCCCGTGTGAAGAGCAAGCAGATCTCCCGCACCTGGGGGGGGACAGGGGG
GCATGGGGACACTGGGGGGACATGGGGGGGGGGGGGGAGGGGGGGGGGG
GGGAGGGGCATGAGGACATTGAGGAGAGGGAACACGAGGGTGGCACTGCA
TCATGGGAGGTGACGAGGGGGTGGGGGGGGCTCAAGGACATGGAGGGGGA
CACTCA

FIGURE 10
SUITE 70

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TTGCTGCCTGCAGGTCGATCTAGTGGATCCGCCGCGACAGCGAACAGGCC
AGCCAGCTGGTGCAGTATCTTTCCACTTTTTTCCGCAAAAACCTTAAAGCG
GCCTTCGGAGTTTGTACTCTCGCCGACGAAATTGAACATGTGAATGCTT
ATCTGCAAATTGAAAAGGCGCGCTTCCAGTCGCGGTTGCAGGTCAACATT
GCTATTCGCAAGAATTATCCCAGCAGCAATTGCCCGCGTTTACCCTGCA
ACCC

FIGURE 10

SUITE 71

20443160 031402

TGCTGGTGGCGGGGATCTGACTGGAAATGGAACGTTCTGTGGCAAGAG
TGGGAATGTAGGAAGGGGGTGGGAGCATGCAGGGTTGGTGGAGCAGGGGG
TAGTGATCAGTGGTGAGGATTTGGTTTCTTGGTCTGAAATATGGATGGAA
GCTTTGTTGGGAGAGTGAATGACTTTTCAGTGAGGACAGGTGGATGCTTG
GGTGAATGCTTGGTAAGTTGTTGAACGCCTGGATAGTTGGATGGGTGGAC
ATGAACCTTTGTATTACAGCTGCAGCTCCAGCACAGAAGGAACCGCCATCC
CAACCACGCCTGGGTGAGCTGACGGCCTCCACGTGAGCCCCGACTCCGT
CCAGCTGGAATGGAGCGTCCCCGAGGGCTCCTTTGACTCCTTCACGGTGC
AGTACAAGGATGCACAAGGCCAGCCACAGGTGGTGCCCGTGGACGGTGGG
TTGCGCACAGTGACCGTGCCCGGGCTGTGCGCGTCCCGCCGCTACAAGTT
CAACCTGTATGGGGTGTGGGGGCGGAAGCGTCTGGGCCCCATGTCCACTG
ATGCTGTACAGGTGAGCATGCTGTTGTGCTGCATCCATGTCTTTTGGCT
GACGGTTGTGTTGGCATATGGTAGGAACCTTTGAGGCCACTCCTGGTTA
CTGTGGTCTTAATAGAGAGGGGAAGTCTTTCTGTCTTGTGACGTGGGTAG
CCTGGAGAGATGGGAGTATGGAAGATGAGAGGAAGAACGGAATAAGGAAT
GATTGATAATTATTGCAGAACGGATGGAAGGGAGGATGGATGGGCGGTGC
ATGGGTACATTGGTGCTTATAGCAGAGCTGGACGGCTGGTTGTACGTTGG
TTTGGTTGTTGAAGAGATGAAGAGTTGGATGGGCGTGTGCTTTCAGTGTG
AATTCCTCCCCCTGTCTTGCAGCTCCGGCACAGAAGGAACCACCTTCCCA
GCCACTCTTGGGTGAGCTGACAGCGTCCACGTGCGCCCCGACTCCGTCC
AGCTGGAATGGAGCGTCCCCGAGGGCTCCTTTGACTCCTTCACGGTGCAG
TACAAGGATGCACAAGGCCAGCCACAGGTGGTGCCCGTGGACGGTGGGTT
GCGCACAGTGACCGTGCCCGGGCTGTGCGCGTCCCGCCGCTACAAGTTCA
ACCTGTATGGGGTGTGGGGGCGGAAGCGTCTGGGCCCCATGTCCACTGAT
GCTGTACAGGTGAGCATGCTGTGTTCTGCCTCCATGTTCTTTTGCTTTC
AGTGTAGTTGTCTGTGTCAGGAACCTTTCTAGGGCCACTTTTGGTTAATG
TTGCCCTTAATAGTCAAGGAACAATTTGTTCTGTTGAGTGGGAATGCCT
AACGGGATGGGAGTTTGGATGATGAGAGGACAAATCTTATAAGGGATGAT
TGATAATTATTGCGGAACGGATGGAAGGAAGGTTGGATGGATGGAATGGT
GTTTGGATAAATTTGTGCTCAGAGCACAGCTGGAGTGTTGGATGAATGTT
GCTTTGCTTGTGTAATAGATGGATGTTTGGTTGTGTGGTTGCTTCCACTG
AGAATTCCTCCCTCTGTGCTGCAGCAGCAGCTCCAGCACAAAGAGGAGCCA
CCTTCCCCACCACGTCTGGGTGAGCTGACAGCGTCCCATGTGCGGCCCCGA
CTCCGTCCAGCTGGAATGGAGCGTCCCCGAGGGCTCCTTTGACTCCTTCA
CGGTGCAGTACAAGGATGCACAAGGCCAGCCACAGGTGGTGCCCGTGGAC
GGTGGGTGCGCACAGTGACCGTGCCCGGGCTGTGCGCGTCCCGCCGCTA
CAAGTTCAACCTGTATGGGGTGTGGGGGCGGAAGCGTCTGGGCCCCATGT
CCACTGATGCTGTACAGGTGAGGGCAGGAATTGGCACCTGTTGGGCTCT
GGGTTTGAGCAGGTAGAAATGTAACAGTGGCCTGCGCTGGGGATCTGTG
TTTCCCCCTGGCAATTGGGAACAGCTGTTGGGTGCCTTTTTTTGGGAAGGATC
CCTTAATCGCAGCATGAAGTATGAATGGACCAATTGGGTGTGGGTGGAGT
GATGGCTGTTGAGATGAGTTGGTGGCTGCTTGAGTAATTGTCTGTTGGAA
TGGATGGACAGATATGTGAAGGAGTGAAAGGATGGATAAAGTAATTTAGG
AATCGGTGGATGAAGAATGGGTAGGTAGACCCTTGGTGAAGTGGTAGAAT
GGAAGGATTTATGAACAGATATGAGTTAATTCTTGCATCGAAGTAGGTGT
AAGTGTCTATTAGCCTGTTGCACTGAACATGCAGTTGCATAGACAAATGA
GTGGGGGAGAAGTACGGAGTAAATCCCTGCATGAATGGTAGGACAGAAACC
TGAATGCCTGGATGCTGGCAGTGTGAAGAATGGCACTTGGGATAGATGGT
TCGAGTATGGGGTAGATTAAGAGATGGATGGAAGAGGAACAGAGAGAG
GGTGATTGGATGAATGGATGGATGGTTGGATGTGACTGATTGACAGGTAC
CAAGCTTTTTTCTGCACTGTGCTTCTGTGCTGCAGCTGCAGAAGAGAC
GGAGGAGGAACACCGTCCCCAGCCACGCCTAGGAGAGCTGACGGCATCCC
ATGTACAGCCCCAACCTCCGTCCAGCTGGAATGGAGCATCCCTGAGGGCTCC
TTTGACTCCTTTCACGGTGCGAGTACATAGACGTGCAAGGCCAGCGCAGGA
GCTGCACTTCTGAGGAGTCTGAGGAGTCTGAGGAGTCTGAGGAGTCTGCTGC

FIGURE 10

SUITE 72

CATCCACCCCTACAAGTTCAACCTTTACGGGGTGTGGGGGCAGACACGT
CTGGGCCCCATCTCCACTGACACCATCACAGGTGAGGGCCCCCTGCCTGCT
GCTGTGCTCTGGGCCTTGTGCTTGGCACGTGGCAGGAGCTGTGCGATGGG
CTGTGCTGGTGGCGGGGATCTGACTGGAAATGGAACGTTCTGTGGCAAA
GAGTGGGAATGTAGGAAGGGGGTGGGAGCATGCAGGGTTGGTGGAGCAGG
GGGTAGTGATCAGTGGTGAGGATTTGGTTTCTTGGTCTGAAATATGGATG
GAAGCTTTGTTGGGAGAGTGAATGACTTTTCAGTGAGGACAGGTGGATGC
TTGGGTGAATGCTTGGTAAGTTGTTGAACGCCTGGATAGTTGGATGGGTG
GACATGAACCTTTGTATTACAGCTGCAGCTCCAGCACAGAAGGAACCGCCA
TCCCAACCACGCCTGGGTGAGCTGACGGCCTCCCACGTGAGCCCCGACTC
CGTCCAGCTGGAATGGAGCGTCCCCGAGGGCTCCTTTGACTCCTTCACGG
TGCAGTACAAGGATGCACAAGGCCAGCCACAGGTGGTGCCCGTGGACGGT
GGGTTGCGCACAGTGACCGTGCCCGGGCTGTGCGCGTCCCGCCGCTACAA
GTTCAACCTGTATGGGGTGTGGGGGCGGAAGCGTCTGGGCCCCATGTCCA
CTGATGCTGTACAGGTGAGCATGCTGTTGTGCTGCATCCATGTCTTTTG
GCTGACGGTTGTGTTGGCATATGGTAGGAACCTTTGAGGCCCACTCCTGG
TTACTGTGGTCTTAATAGAGAGGGAAGTTCTTTCCTGTTCTTGACGTGGG
TAGCCTGGAGAGATGGGAGTATGGAAGATGAGAGGAAGAACGGAATAAGG
AATGATTGATAATTATTGCAGAACGGATGGAAGGGAGGATGGATGGGCGG
TGCATGGGTACATTGGTGCTTATAGCAGAGCTGGACGGCTGGTTGTACGT
TGGTTTGGTTGTTGAAGAGATGAAGAGTTGGATGGGCGTGTGCTTTCCT
GTGAATTCCTCCCCCTGTCTTGACAGCTCCGGCACAGAAGGAACCACCTTC
CCAGCCACTCTTGGGTGAGCTGACAGCGTCCCACGTGCGCCCCGACTCCG
TCCAGCTGGAATGGAGCGTCCCCGAGGGCTCCTTTGACTCCTTCACGGTG
CAGTACAAGGATGCACAAGGCCAGCCACAGGTGGTGCCCGTGGACGGTGG
GTTGCGCACAGTGACCGTGCCCGGGCTGTGCGCGTCCCGCCGCTACAAGT
TCAACCTGTATGGGGTGTGGGGGCGGAAGCGTCTGGGCCCCATGTCCACT
GATGCTGTACAGGTGAGGGCAGGAATTGGCACCTGGTGGGCTCTGGGT
TGCAGCAGGTAGAAATGTAAACGTGGCCTGCGCTGGGGATCTTGTTTTCC
CCTGGCAATGGGAACAGCTGTTGGGTGCCTTTTTTGGGAAGGATCCCTTA
ATCGCAGCATGAAGTATGAATGGACCAATTGGGTGTGGGTGGAGTGATGG
CTGTTGAGATGAGTTGGTGGCTGCTTGAGTAATTGTCTGTTGGAATGGAT
GGACAGATATGTGAAGGAGTGAAGGATGGATAAAGTAATTTAGGAATCG
GTGGATGAAGAATGGGTAGGTAGACCCTTGGTGAAGTGGTAGAATGGAAG
GATTTATGAACAGATATGAGTTAATTCTTGATCGAAGTAGGTGTAAGTG
TCTATTAGCCTGTTGCACTGAACATGCAGTTGCATAGACAAATGAGTGGG
GAGAAGTACGGAGTAAATCCCTGCATGAATGGTAGGACAGAAACCTGAAT
GCCTGGATGCTGGCAGTGTGAAGAATGGCACTTGGGATAGATGGTTCGAG
TATGGGGTAGATTAAAAGATGGATGGAAAAGAGGAACAGAGAGAGGGTGA
TTGGATGAATGGATGGATGGTTGGATGTGACTGATTGACAGGTACCAAGC
TTTTTTCCTGCACTGTGCCTTCTGTGCTGCAGGACTATGGTCATAGCTGT
TTCCTGTGTGAAATTGTTATCCGCTCACAATTCCACACAACATCGA

FIGURE 10

SUITE 73

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